

Microarrays

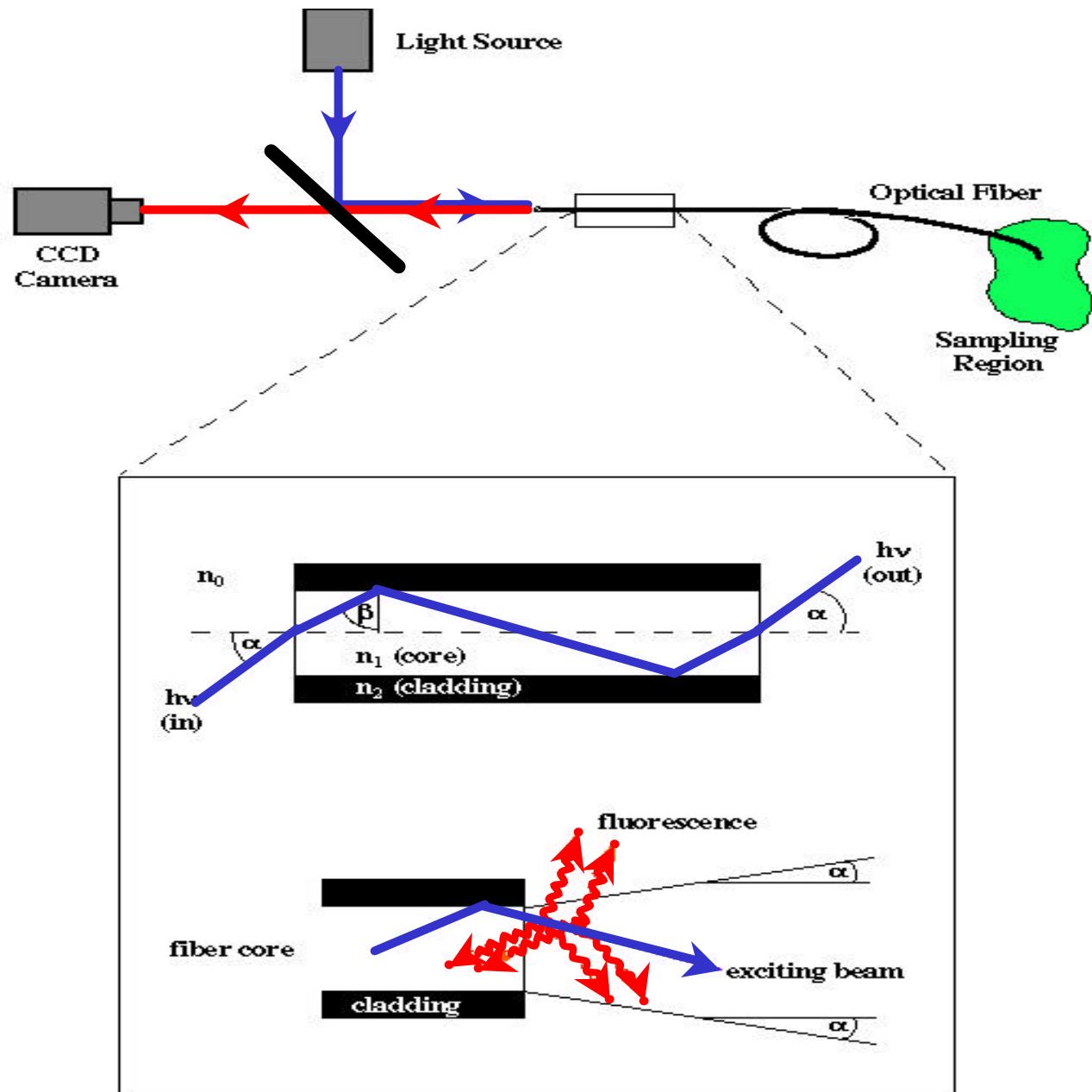
New England Bioterrorism Preparedness Workshop

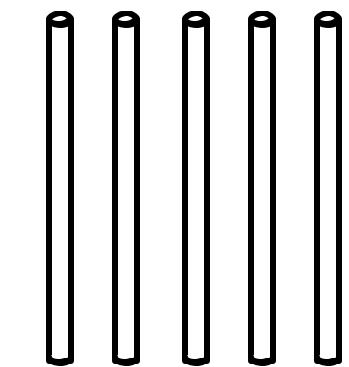
Dr. David Walt

Tufts University

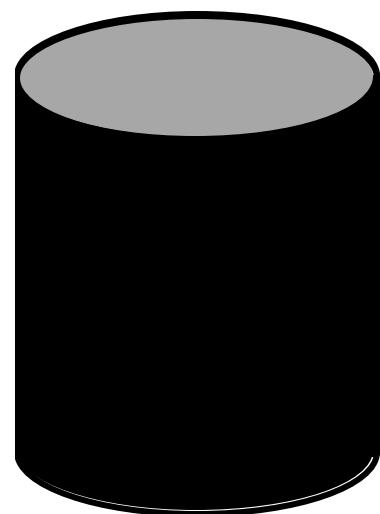
4 April 2002

Report Documentation Page		
Report Date 04APR2002	Report Type N/A	Dates Covered (from... to) 03APR2002 - 04APR2002
Title and Subtitle Microarrays		Contract Number F19628-00-C-0002
		Grant Number
		Program Element Number
Author(s) Walt, David		Project Number
		Task Number
		Work Unit Number
Performing Organization Name(s) and Address(es) Tufts University		Performing Organization Report Number
Sponsoring/Monitoring Agency Name(s) and Address(es) Air Force ESC/XPK (Richard Axtell) Hanscom AFB, MA 01731		Sponsor/Monitor's Acronym(s)
		Sponsor/Monitor's Report Number(s)
Distribution/Availability Statement Approved for public release, distribution unlimited		
Supplementary Notes Workshop paper from the New England Bioterrorism Preparedness Workshop held 3-4 april 2002 at MIT Lincoln Laboratory, Lexington, MA, The original document contains color images.		
Abstract		
Subject Terms		
Report Classification unclassified		Classification of this page unclassified
Classification of Abstract unclassified		Limitation of Abstract SAR
Number of Pages 69		

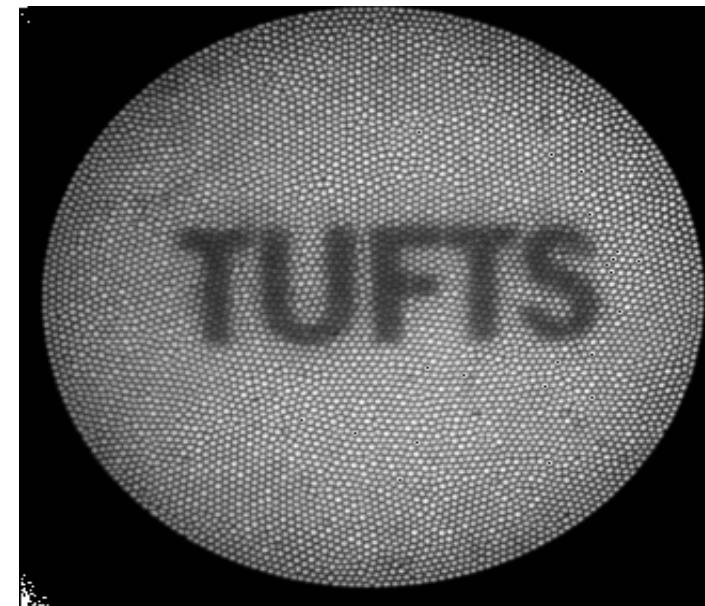




Individual Cladded
Optical Fibers



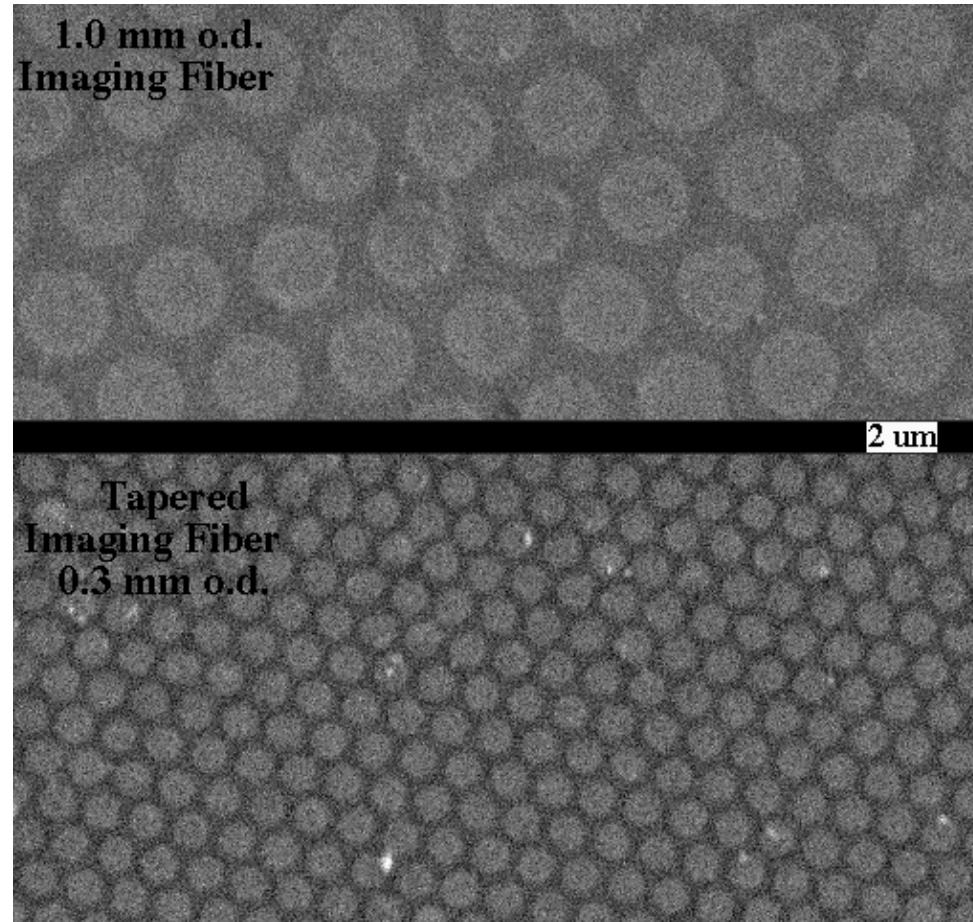
Silica
Jacket



Imaging
Fiber



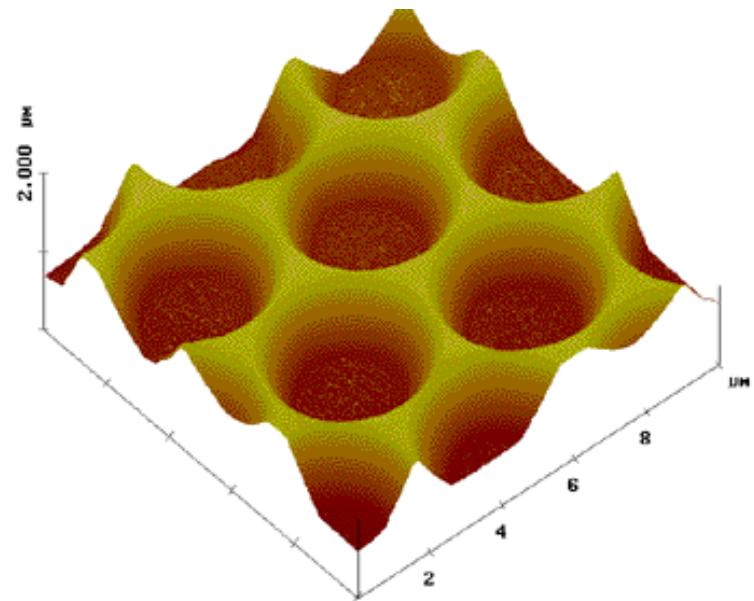
Optical Imaging Fiber Before and After Tapering



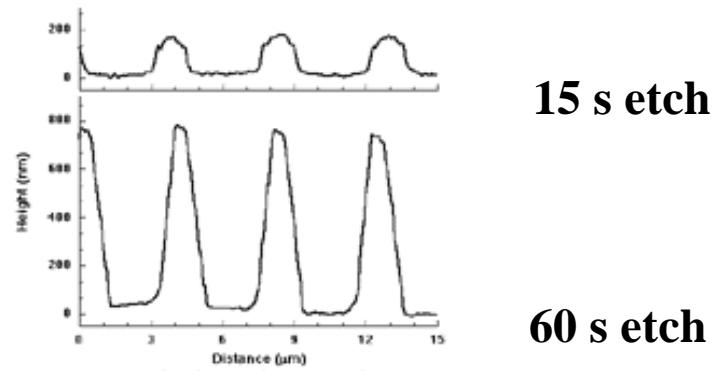
**Individual Core
Diameter $\sim 2.6 \text{ } \mu\text{m}$**

**Individual Core
Diameter $\sim 0.85 \text{ } \mu\text{m}$**

AFM of a Chemically-Etched 1000-?m Diameter Imaging Fiber

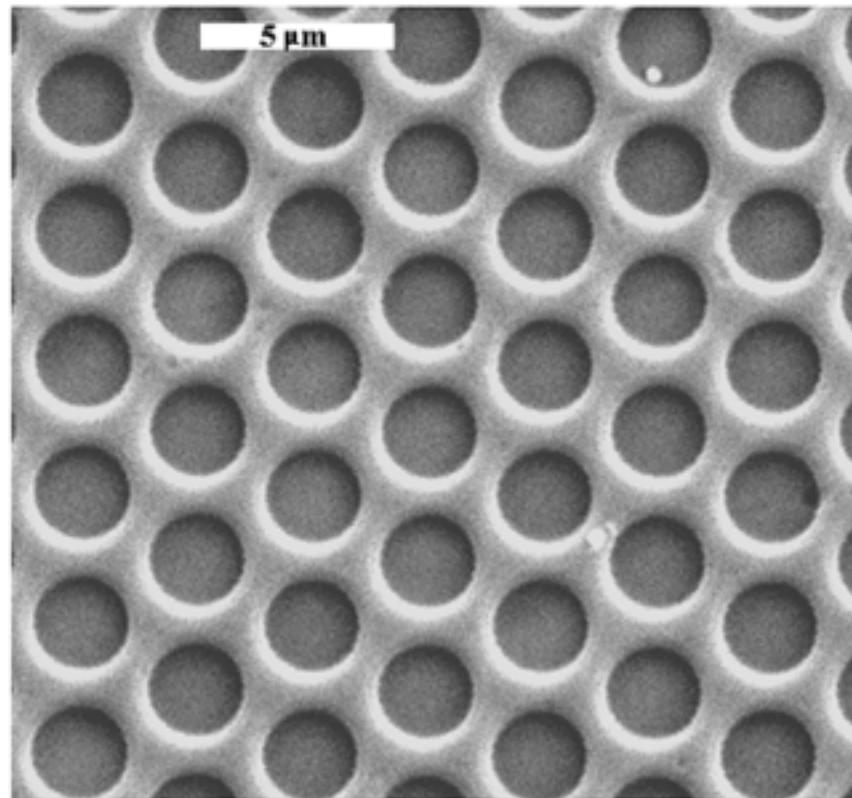


Well Profiles



Pantano, P.; Walt, D.R. *Chem. Mater.* **1996**, 8, 2832-2835

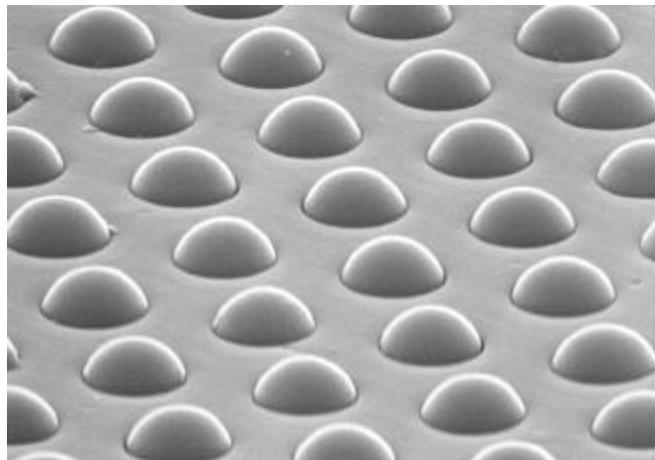
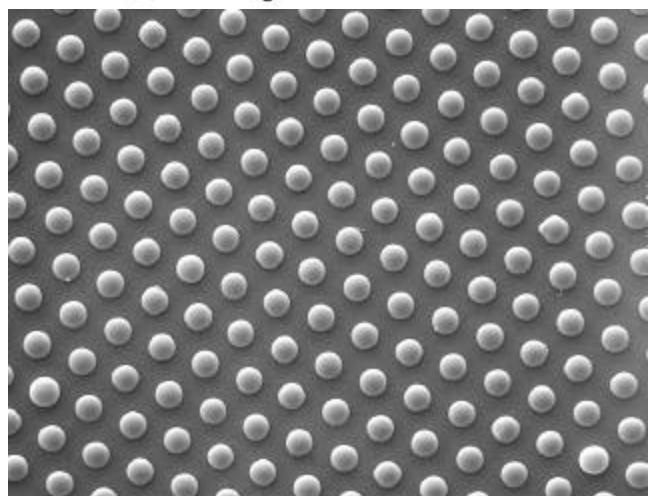
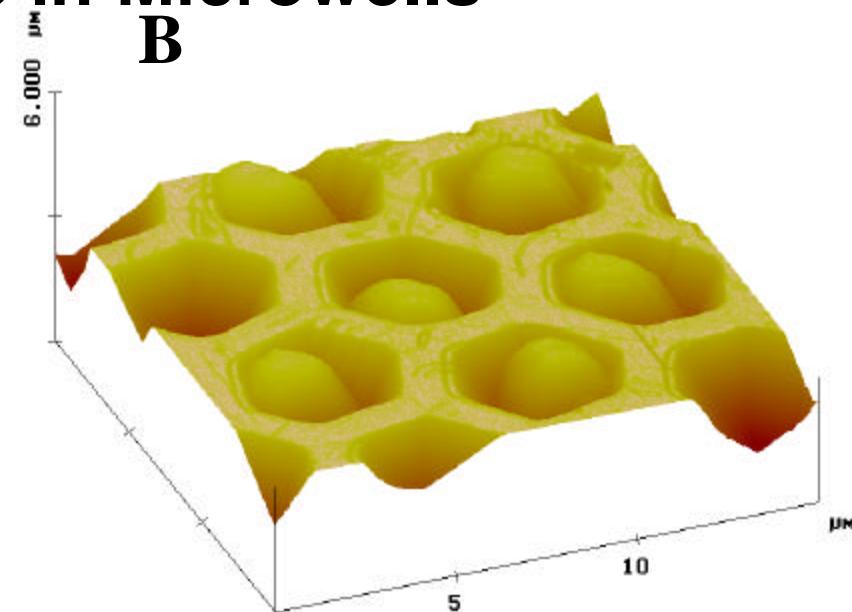
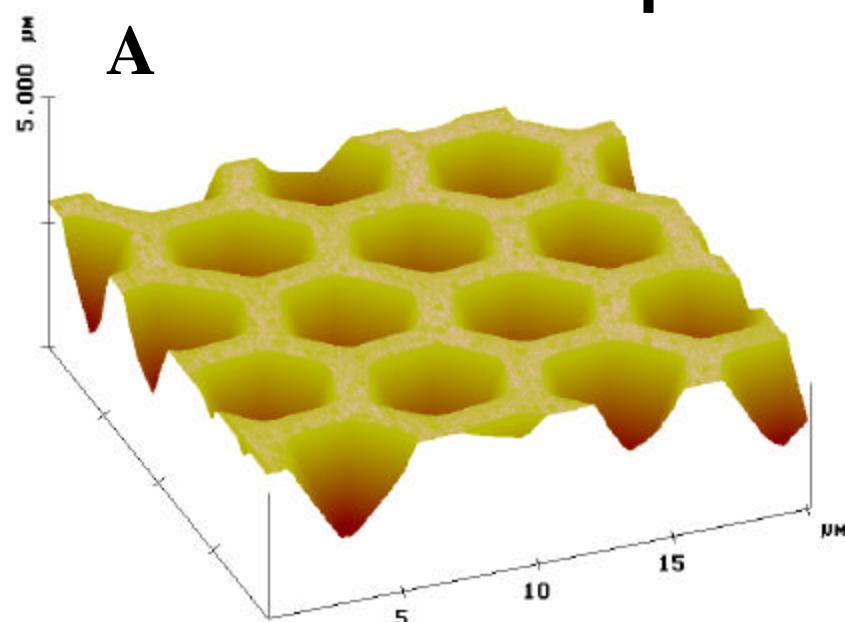
SEM of a Chemically-Etched 1000-? m Diameter Imaging Fiber



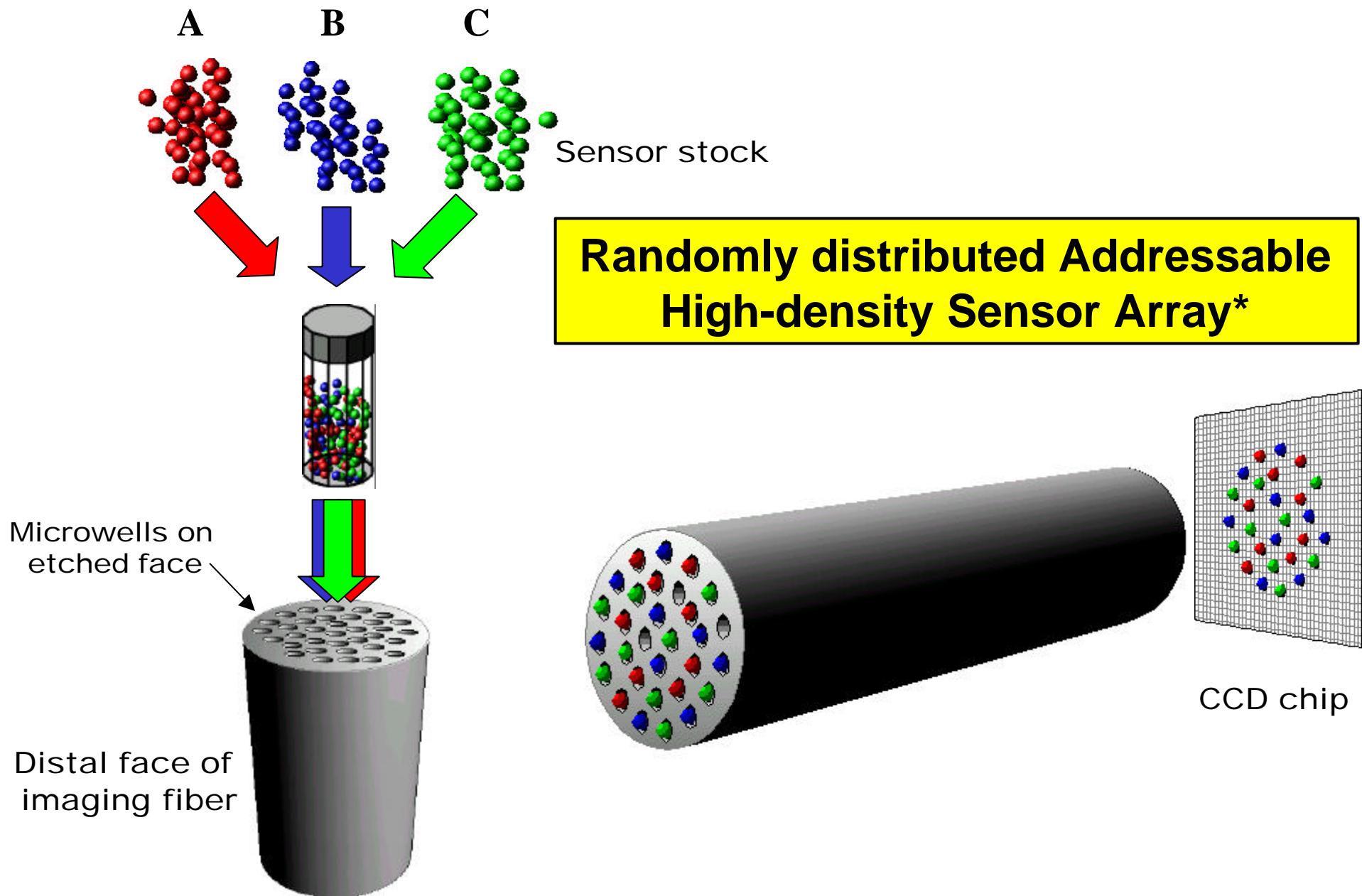
Pantano, P.; Walt, D.R. *Chem. Mater.* **1996**, 8, 2832-2835

Microspheres in Microwells

B

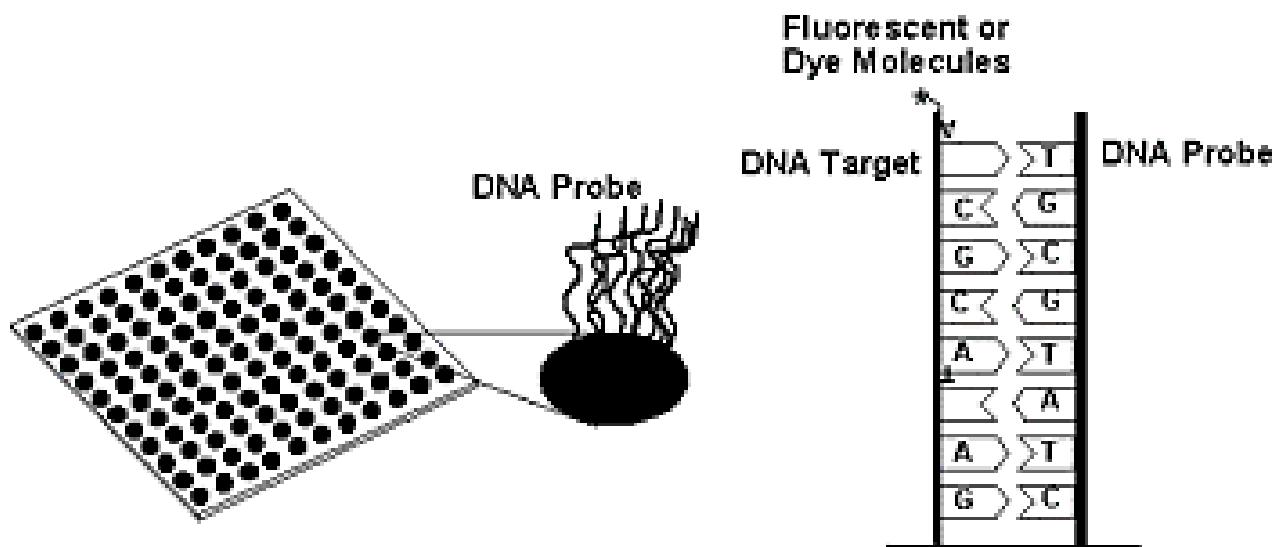


Michael, K.L. et al. *Anal. Chem.* 70 (7): 1242-1248 (1998).

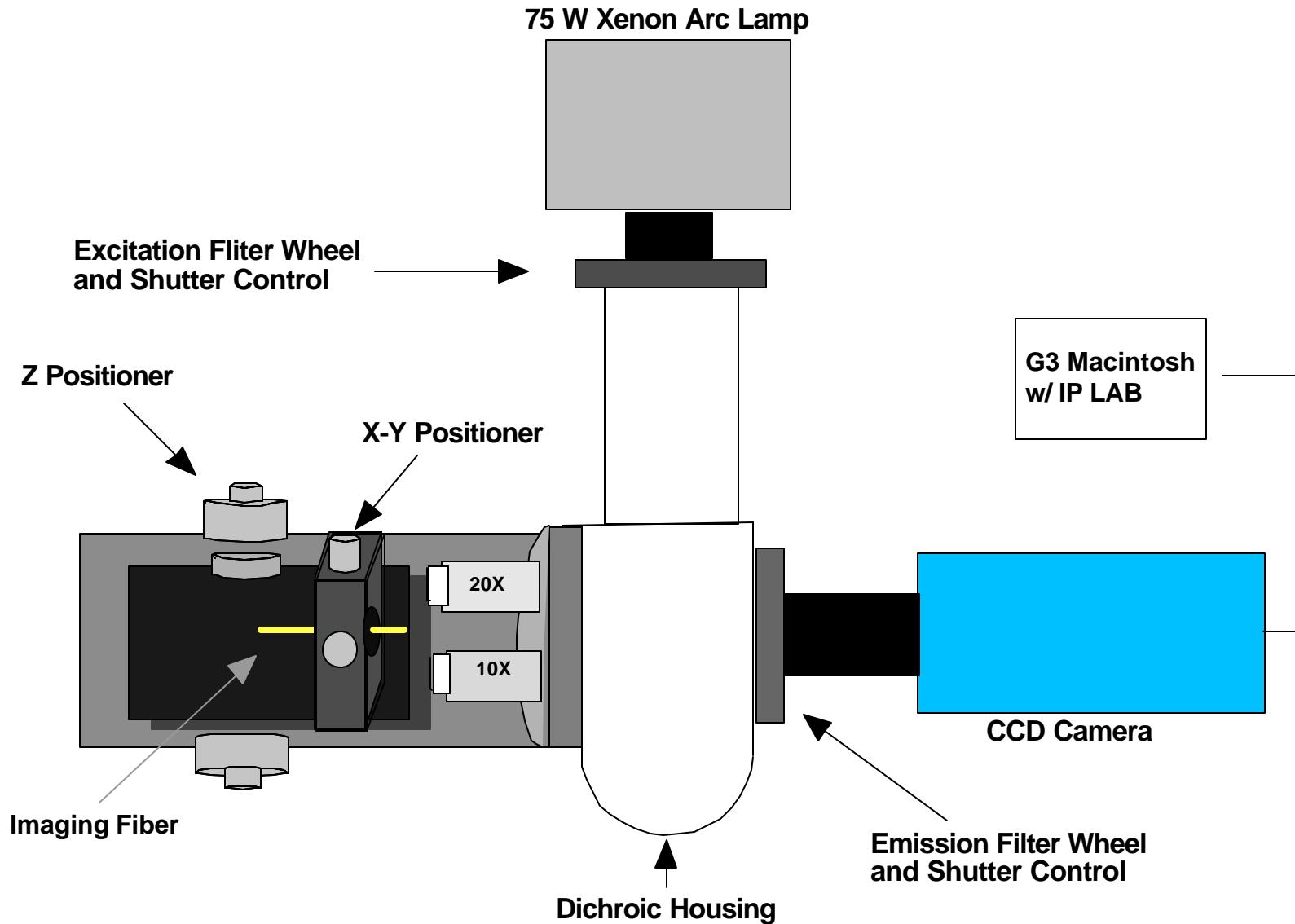


*Michael et al. 1998 *Anal. Chem.* **70**: 1242-1248

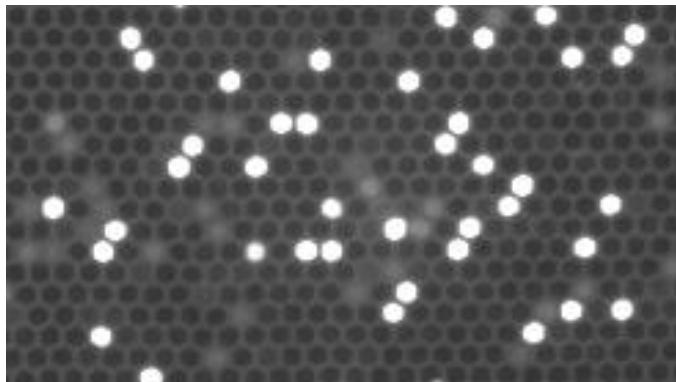
DNA Array Principle



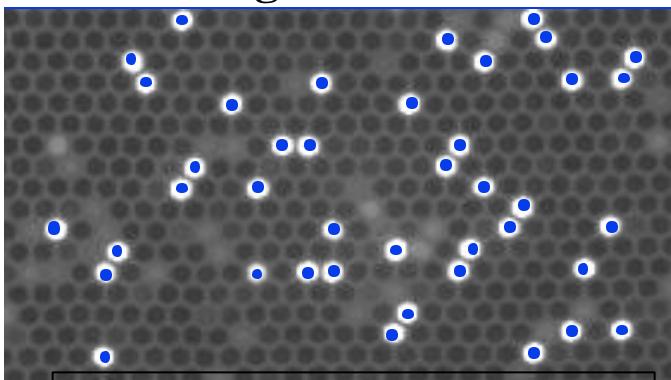
Instrumentation: Modified Fluorescence Microscope



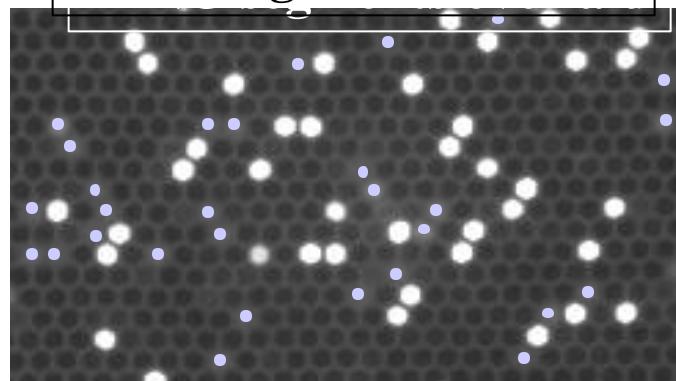
Encoding Signal of Dye 1



Signal 530

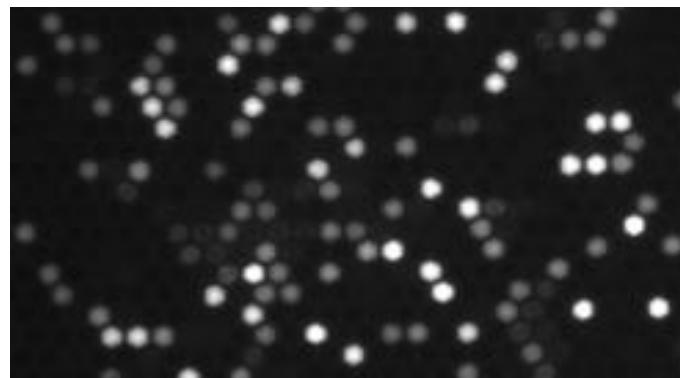


IFNG segments overlaid

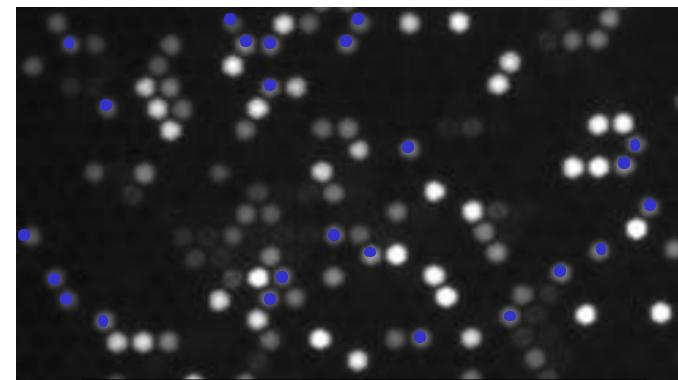


IL2 segments overlaid

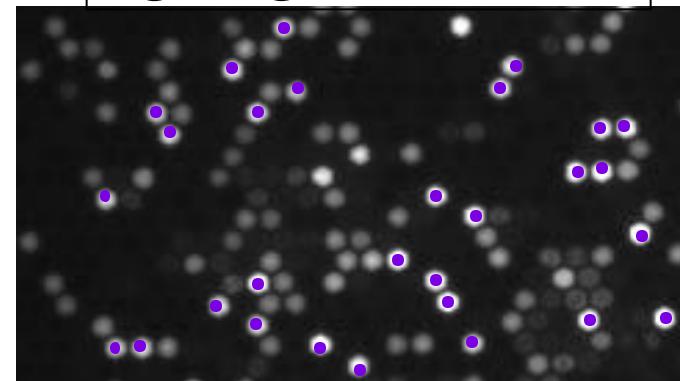
Encoding Signal of Dye 2



Signal 620



Bglo segments overlaid



Hwt segments overlaid

Sequences of 25 Probes used together in a Microsphere Array

1) ?-glo (segment of human ?-globin)²⁶
TCA ACT TCA TCC ACG TTC ACC

2) IFNG (interferon gamma 1)²⁶
IFNG TGG GTT CTC TTG GCT GTT ACT

3) IL2 (interleukin-2)²⁶
TA CAA GAA TCC CAA ACT CAC CAG

4) IL4 (interleukin-4)²⁶
CC AAC TGC TTC CCC CTC TGT

5) IL6 (interleukin-6)²⁶
GT TGG GTC AGG GGT GGT TAT T

6) K-ras WT²⁷
GGA GCT GGT GGC GTA

7) H-ras WT²⁷
CCG GCG GTG T

8) CFTR (cystic fibrosis exon 11)¹³
CAT TAT ACT TGT AGA G

9) R553X (cystic fibrosis exon 10)¹³
TGT AGA ATT ATC TTC

10) PAN132¹⁶ (human peripheral lymphocyte)
CCT CTA TAC TTT AAC GTC AAG

11) Schena-2¹⁶
AAG TTT AAC CTA TAC CCT GTC

12) Hakala-1²⁰
CCT ATG ATG AAT ATA G

13) Hakala-2²⁰
AAT ATG ATA ATG GCC T

14) complement to probe 1
TG AAC GTG GAT GAA GTT G

15) complement to probe 2
AG TAA CAG CCA AGA GAA CCC AAA

16) complement to probe 3
CT GGT GAG TTT GGG ATT CTT GTA

17) complement to probe 4
AC AGA GGG GGA AGC AGT TGG

18) complement to probe 5
AA TAA CCA CCC CTG ACC CAA C

19) complement to probe 6
TAC GCC ACC AGC TCC

20) complement to probe 7
ACA CCG CCG G

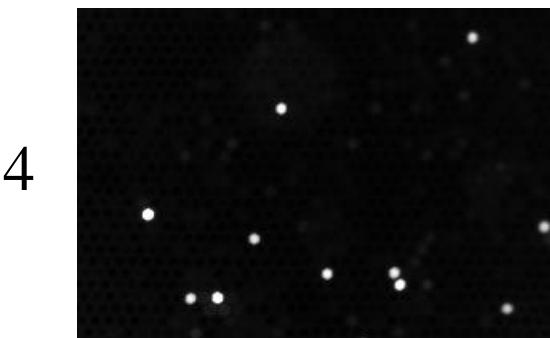
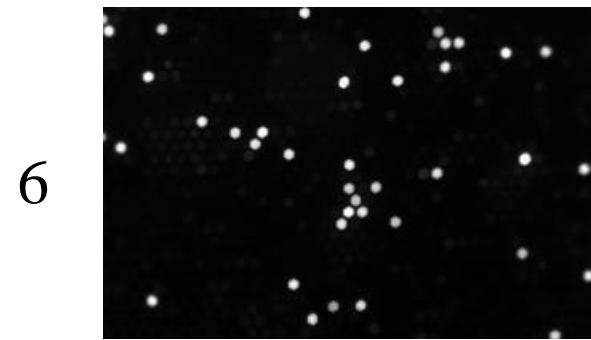
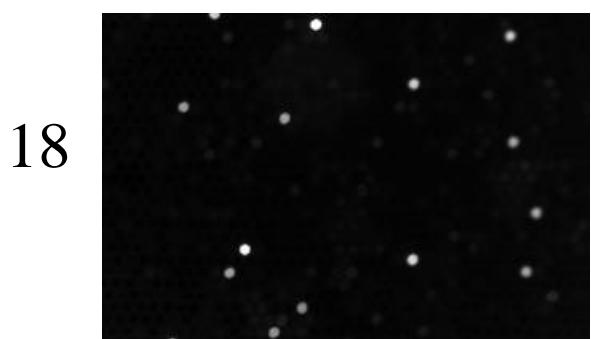
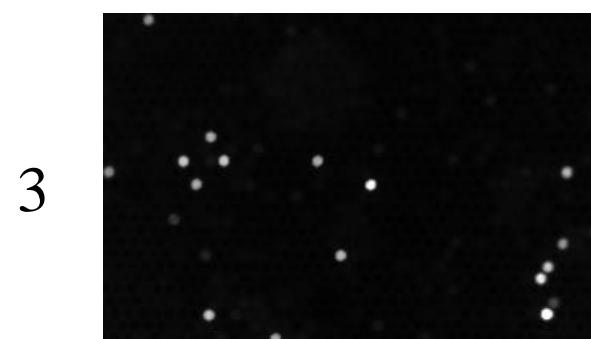
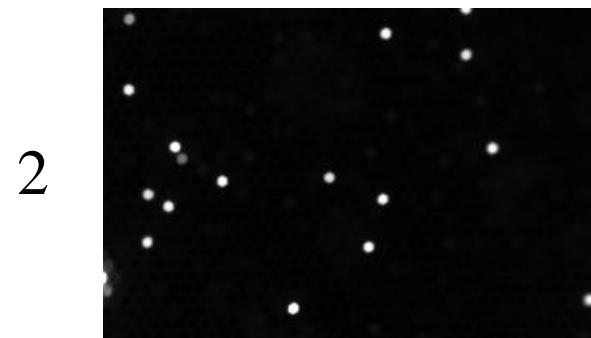
21) complement to probe 8
CTC TAC AAG TAT AAT G

22) complement to probe 9
GAA GAT GTT AAA GTA TAG AGG

23) complement to probe 10
CTA GAC GTT AAA GTA TAG AGG

24) complement to probe 12
CTA TAT TCA TCA TAG G

25) complement to probe 13
AGG CCA TTA TCA TAT T



E. coli Allelic Discrimination

ycgW locus*

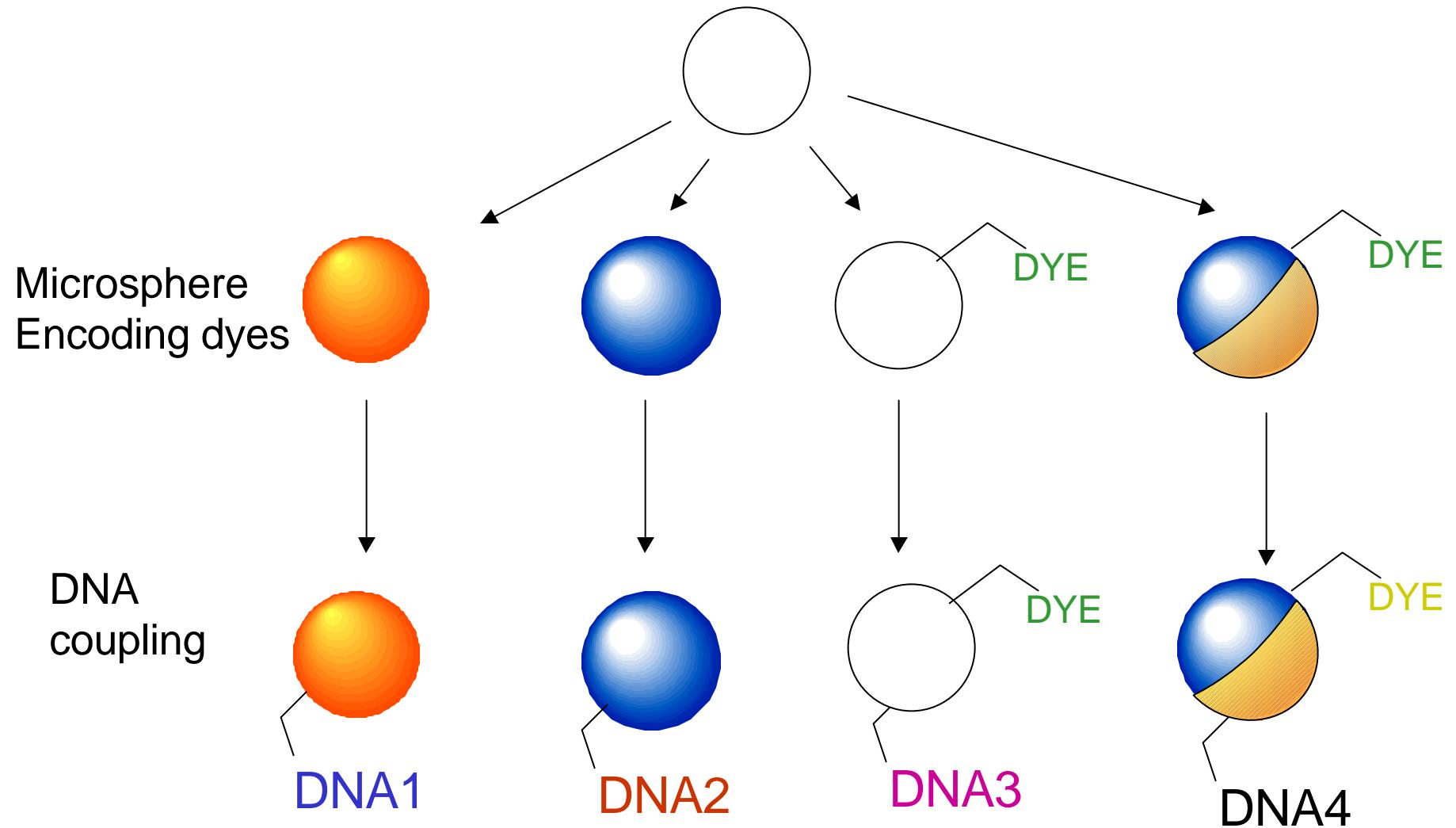
No.	strain	strain sequence												No.	strain	strain sequence											
1	BSR9b	T	T	T	T	T	G	A	A	G	G	G	G	19	HER1265	no PCR product											
2	BSR9c	T	T	T	T	T	G	A	A	G	G	G	G	20	HER1266	no PCR product											
3	"ETEC"	T	T	T	T	T	T	G	A	G	G	G	G	21	EC68	no PCR product											
4	O111NM	T	T	T	T	T	G	A	A	G	G	G	G	22	EC69	T	T	T	T	T	G	A	G	G	G	G	G
5	O113:H2	T	T	T	T	T	G	A	A	G	G	G	G	23	EC63	no PCR product											
6	O157NM	T	T	T	T	T	T	G	A	G	G	G	G	24	EC54	no PCR product											
7	HER1058	no PCR product												27	O86:H10	T	T	T	T	T	G	A	A	G	G	G	
8	K12DH5a	T	T	T	T	T	T	G	A	G	G	G	G	37	O86:H18	T	T	T	G	T	T	T	T	T	T	T	G
9	K12W4100	T	T	T	T	T	T	G	A	G	G	G	G	30	O8:H9	T	T	T	T	T	G	A	A	G	G	G	G
10	O55:H7	T	T	T	T	C	G	A	A	G	G	G	G	34	O9:H33	T	T	T	T	T	G	A	G	G	G	A	G
11	"EPEC"	T	T	T	T	T	T	G	A	G	G	G	G	38	O153:H-	T	T	T	T	T	G	A	G	G	G	G	G
12	K12W3110	T	T	T	T	T	T	G	A	G	G	G	G	43	O26:H11	T	T	T	T	T	G	A	A	G	G	G	G
13	O22:H8	T	T	T	T	T	G	A	A	G	G	G	G	48	O127:H21	T	T	T	T	T	G	A	A	G	G	G	G
14	O26:H-	T	T	T	T	T	T	G	A	G	G	G	G	52	EC1	T	T	T	T	T	G	A	G	G	G	G	G
15	O42:H2	T	T	T	T	T	G	A	A	G	G	G	G	53	EC7	T	T	T	T	T	G	A	G	G	G	G	G
16	O157:H7	no PCR product												54	EC18	T	T	T	T	T	G	A	G	G	G	G	
17	HER1057	no PCR product												55	EC47	T	T	T	G	T	T	T	T	T	T	T	G
18	HER1261	no PCR product												56	EC52	no PCR product											
	Cons.	T	T	T	*	*	*	*	*	*	*	*	*	Cons.	T	T	T	*	*	*	*	*	*	*	*	*	

= probe 1 signal

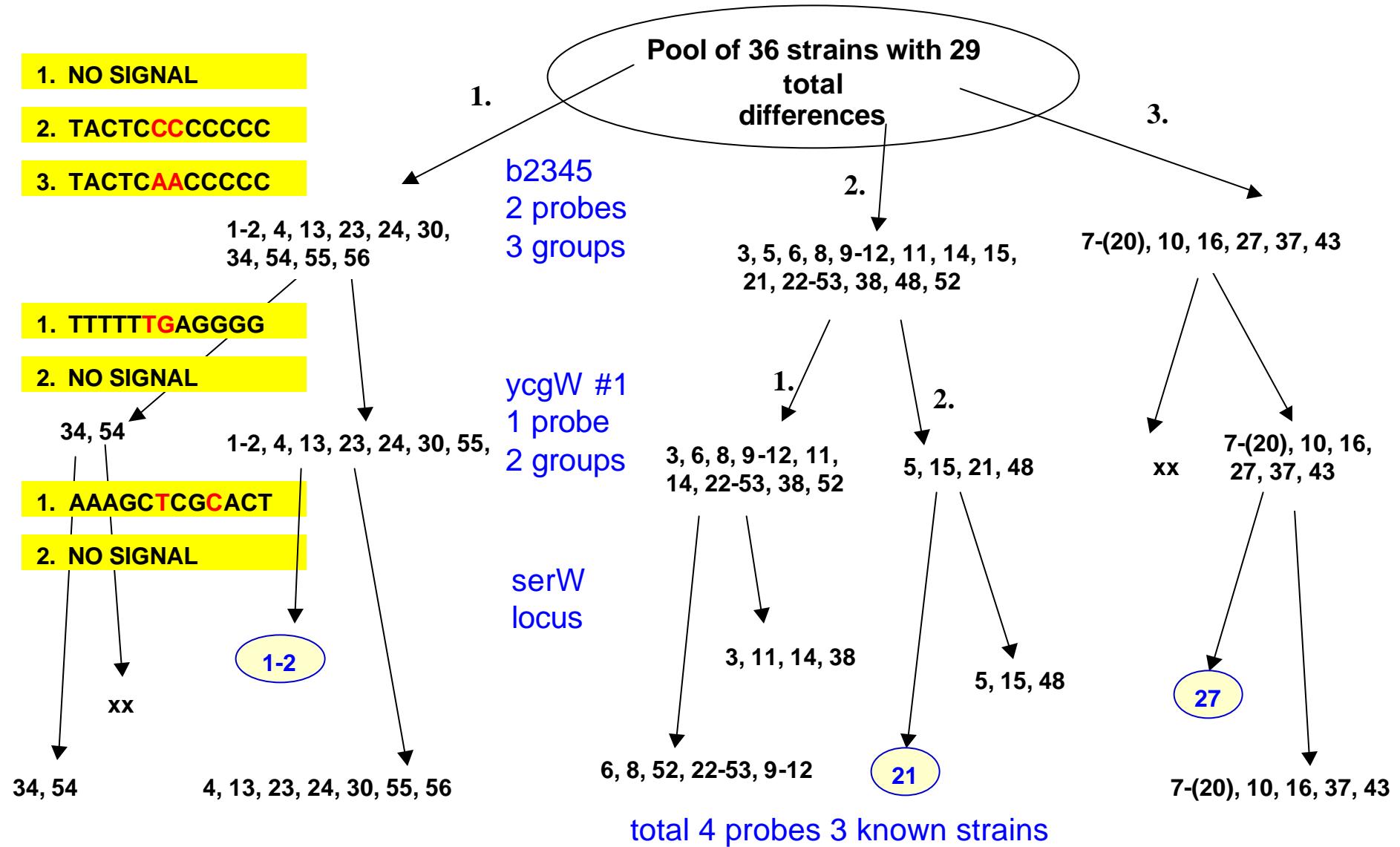
= no signal

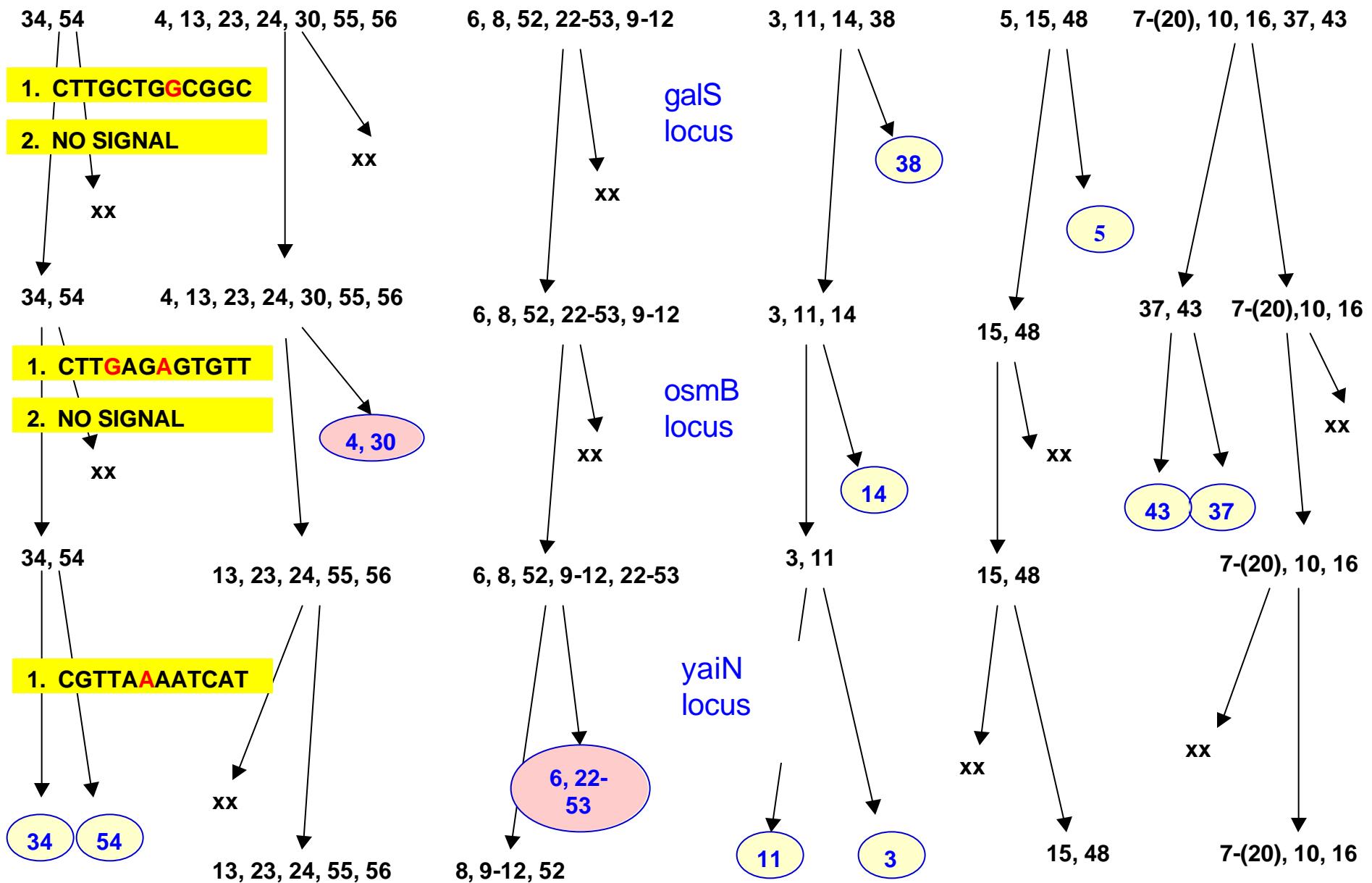
*ycgW locus is 77 nucleotides long

Microsphere Functionalization

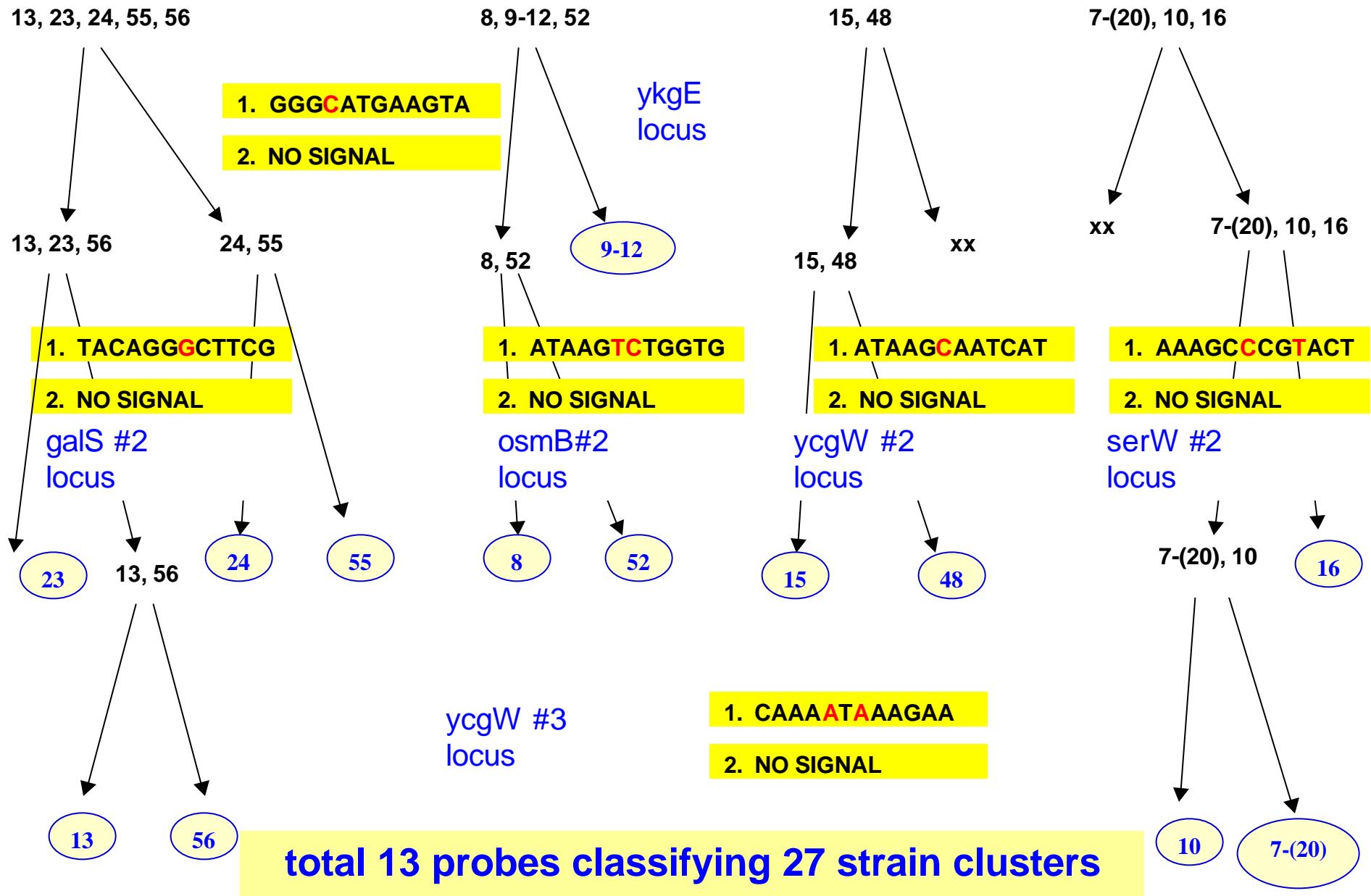


E. coli Genomic Discrimination Flowchart





total 7 probes 12+ known strains



E. coli Genomic Pattern Response

No.	strain	ycgW	serW	osmB	yaiN	ykgE
14	O26:H-	blue	yellow	yellow		
11	"EPEC"	blue	yellow	blue	blue	
3	"ETEC"	blue	yellow	blue	yellow	
6	O157NM	blue	blue	blue	yellow	
22	EC69	blue	blue	blue	yellow	
9	K12W4100	blue	blue	blue	blue	yellow
12	K12W3110	blue	blue	blue	blue	yellow
8	K12DH5a	blue	blue	blue	blue	blue



= signal response



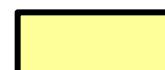
= no signal

E. coli Genomic Pattern Response

No.	strain	ycgW	b2345	serW	galS	Osmb	yKgE	ycgW#3	serW#2
21	EC68	Yellow	Blue	Blue					
1	BSR9b	Yellow	Yellow	Blue					
2	BSR9c	Yellow	Yellow	Blue					
15	O42:H2	Yellow	Blue	Yellow	Blue				
5	O113:H2	Yellow	Blue	Yellow	Yellow				
4	O111NM	Yellow	Yellow	Yellow	Blue	Yellow			
24	EC54	Yellow	Yellow	Yellow	Blue	Blue	Yellow		
13	O22:H8	Yellow	Yellow	Yellow	Blue	Blue	Blue		
23	EC63	Yellow	Yellow	Yellow	Blue	Blue	Yellow		
10	O55:H7	Yellow	Yellow	Yellow	Yellow	Blue	Yellow	Blue	
16	O157:H7	Yellow	Yellow	Yellow	Yellow	Blue	Yellow	Yellow	Yellow
7	HER1058	Yellow	Yellow	Yellow	Yellow	Blue	Yellow	Yellow	Blue
17	HER1057	Yellow	Yellow	Yellow	Yellow	Blue	Yellow	Yellow	Blue
18	HER1261	Yellow	Yellow	Yellow	Yellow	Blue	Yellow	Yellow	Blue
19	HER1265	Yellow	Yellow	Yellow	Yellow	Blue	Yellow	Yellow	Blue
20	HER1266	Yellow	Yellow	Yellow	Yellow	Blue	Yellow	Yellow	Blue

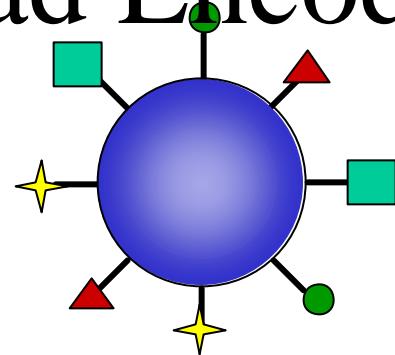


= signal response

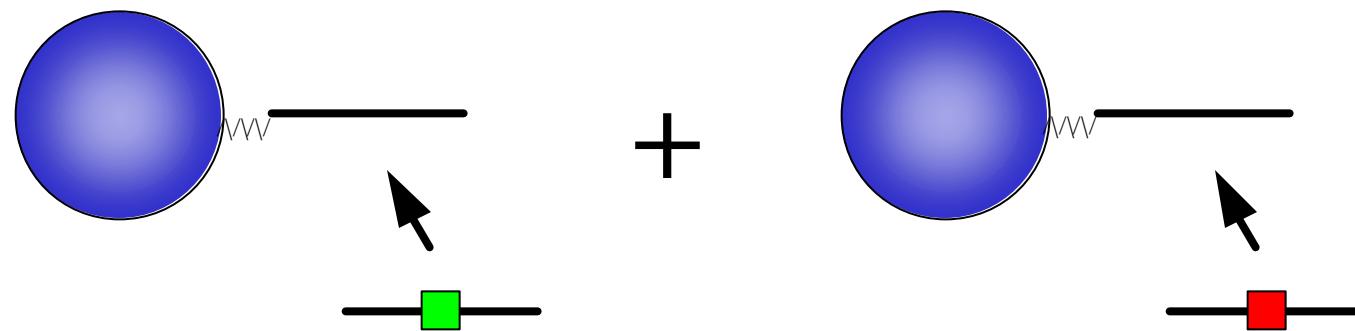


= no signal

Bead Encoding

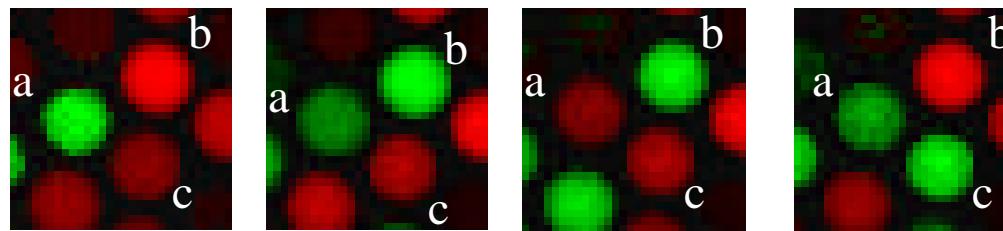


Sequential Decoding



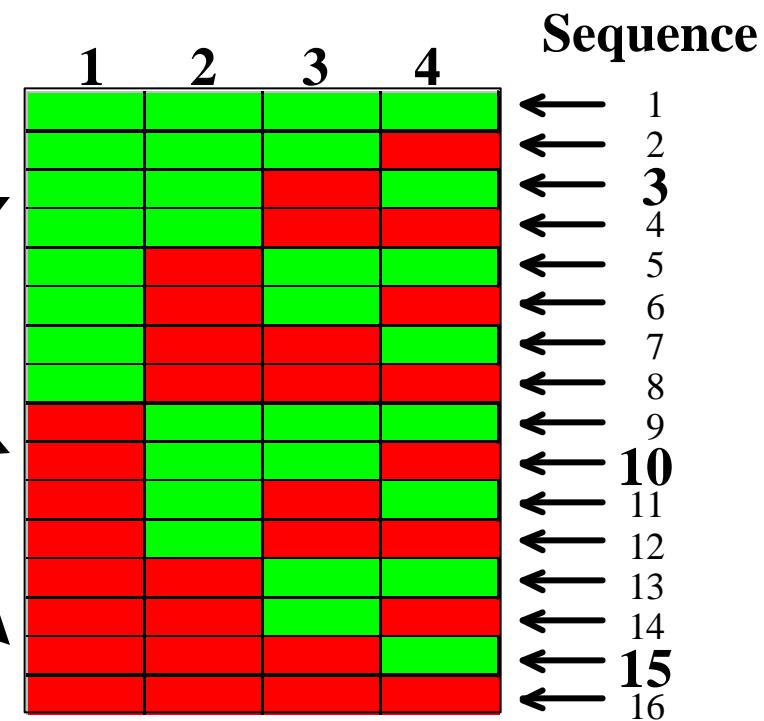
Decoding 16 Probes

Hyb # 1 2 3 4



Hyb.#1 Hyb.#2 Hyb.#3 Hyb.#4

a	green	green	red	green
b	red	green	green	red
c	red	red	red	green

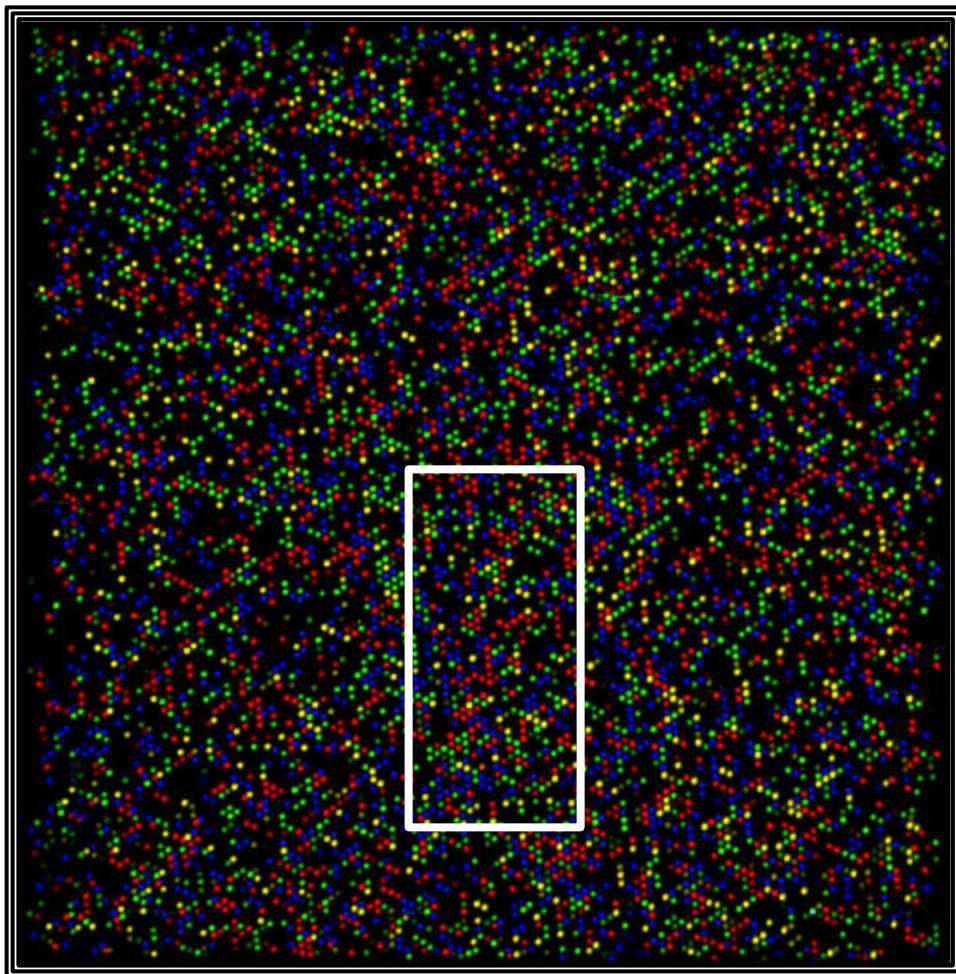


Decoding is Exponential

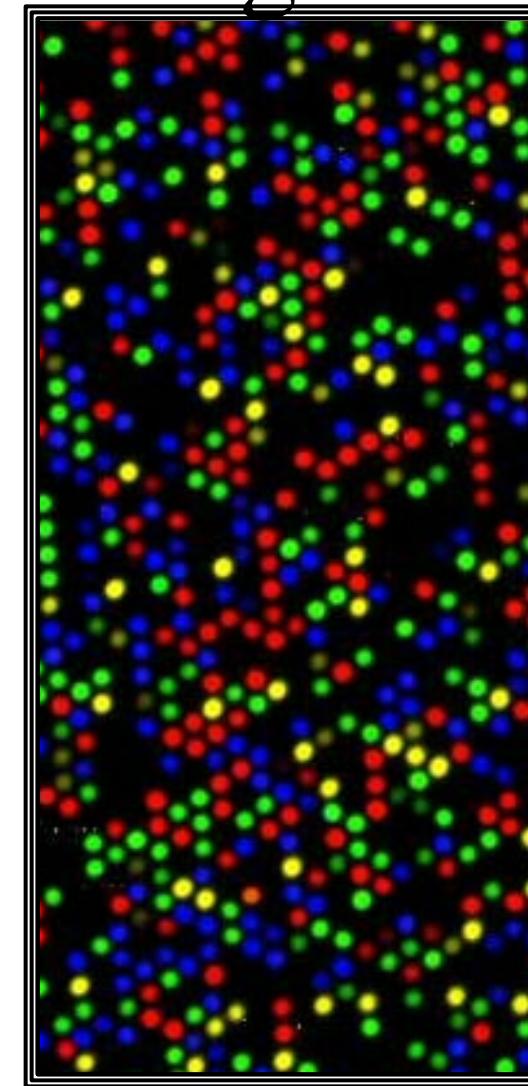
x labels , z steps = x^z codes

- 2 Dyes \wedge 4 Steps = 16 Codes
- 4 Dyes \wedge 6 Steps = 4,096 Codes

Four-Color Decoding

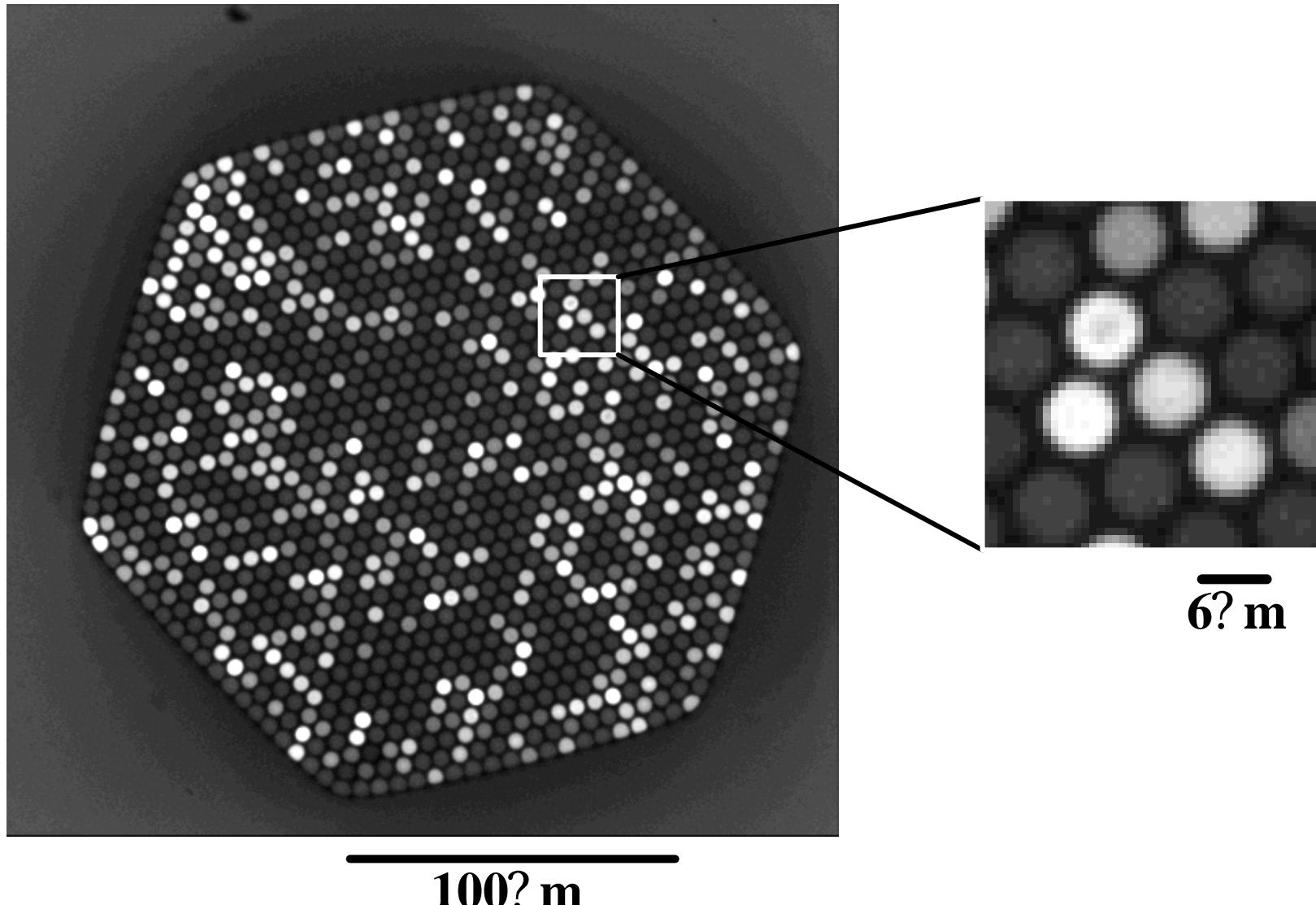


~13,000 Wells, 16 Probe Sequences

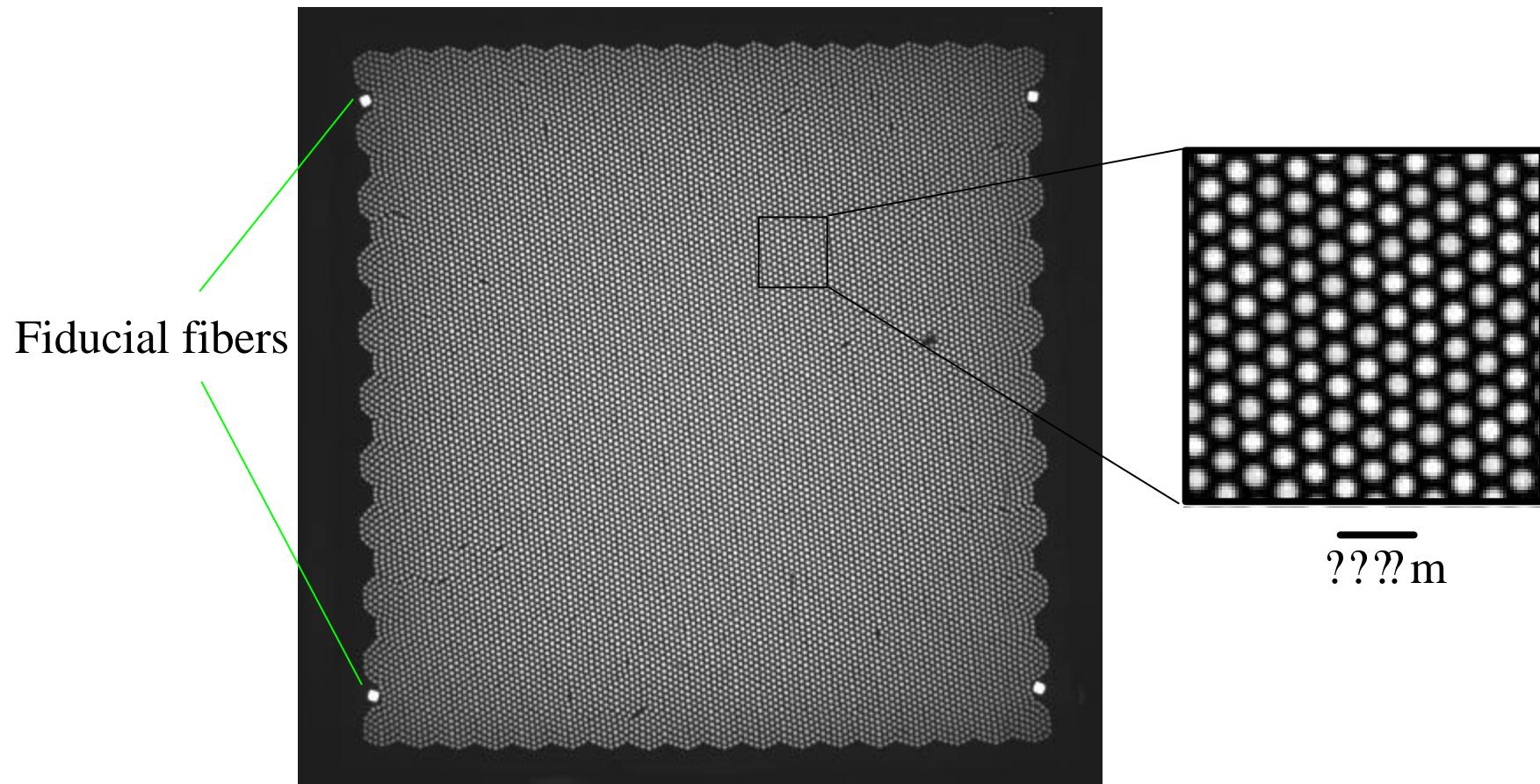


D.R. Walt, *Science*, 2000

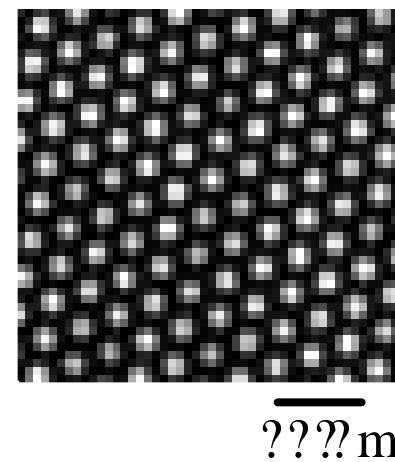
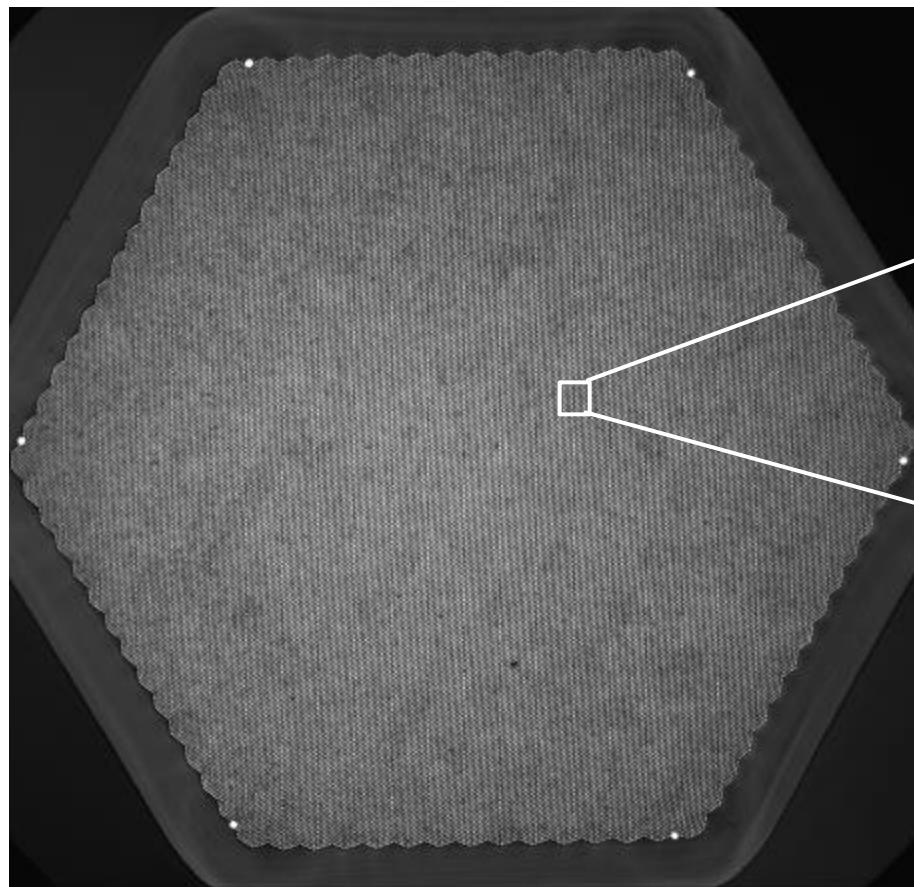
1K Fiber Bundle



13K Fiber Bundle

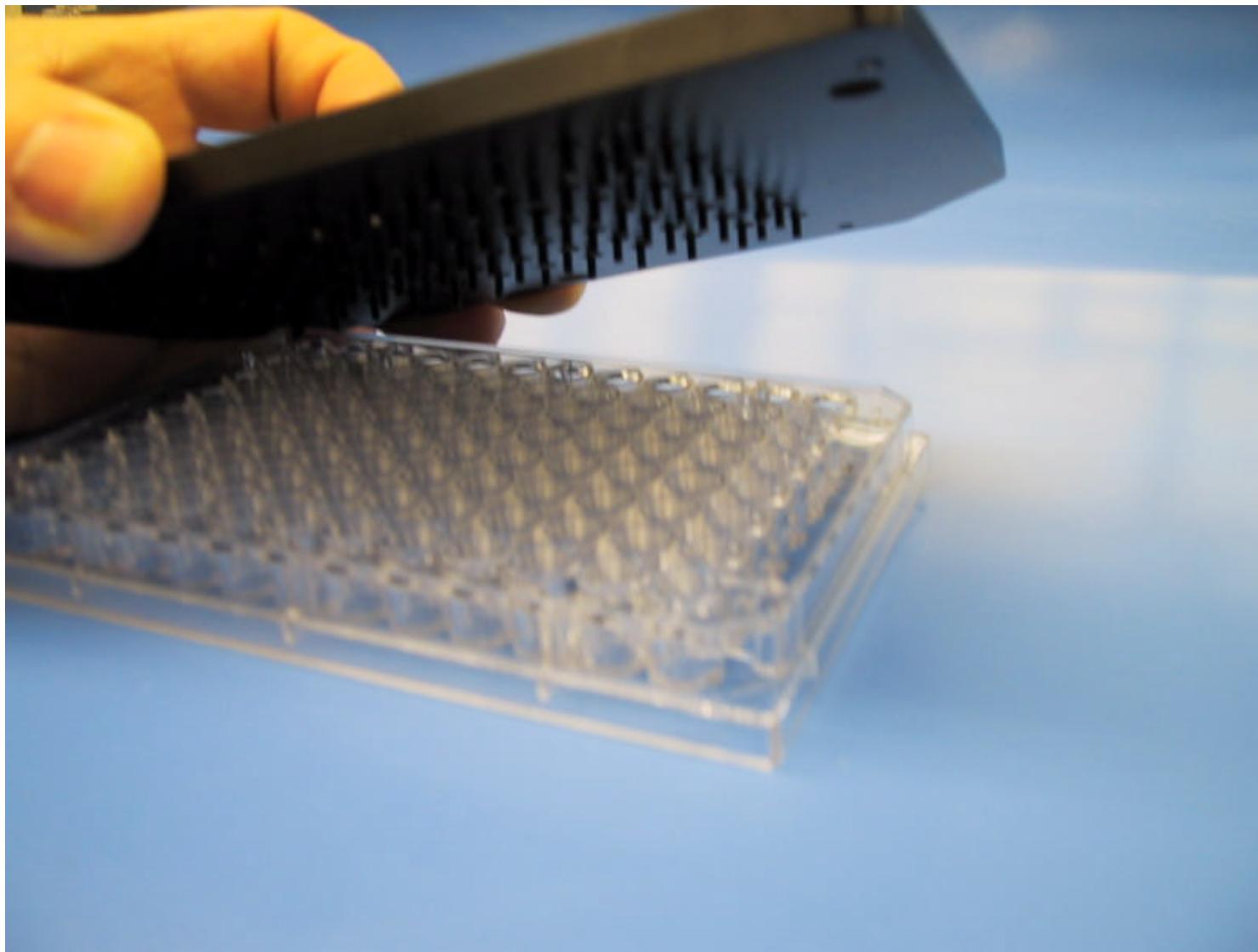


57K Fiber Bundle



??? m

Array of Arrays™



Scalability of Technology



Unique Experiments
(with ~20-fold redundancy)



Bead Summing

Concentration	Hybridization Time: No Summing	Hybridization Time: 100 Bead Summed
100 pM	10 minutes	4 minutes
10 pM	30 minutes	7 minutes
100 fM	4 hours	20 minutes
10 fM	17 hours	30 minutes

Size and Concentration

Volume			1 ?? M	1 nM	1 pM
$(1 \text{ mm})^3$	1 L	10^{-6} L	6×10^{-11}	6×10^{-8}	6×10^{-5}
$(100 \text{ ? m})^3$	1 nL	10^{-9} L	6×10^{-8}	6×10^{-5}	6×10^{-2}
$(10 \text{ ? m})^3$	1 pL	10^{-12} L	6×10^{-5}	6×10^{-2}	6×10^{-1}
$(1 \text{ ? m})^3$	1 fL	10^{-15} L	6×10^{-2}	6×10^{-1}	
$(0.1 \text{ ? m})^3$	1 aL	10^{-18} L	6×10^{-1}		

Probe and Target Sequences for DNA Microarray Detection Limits

Probe

IL2 (interleuken-2) 5'-TA-CAA-GAA-TCC-CAA-ACT-CAC-CAG-3'

IL6 (interleuken-6) 5'-GT-TGG-GTC-AGG-GGT-GGT-TAT-T-3'

F508C 5'-TAG-GAA-ACA-CCA-CAG-ATG-ATA-3'

Target

IL2 (interleuken-2) 5'-CT-GGT-GAG-TTT-GGG-ATT-CTT-GTA-3'

IL6 (interleuken-6) 5'-AA-TAA-CCA-CCC-CTG-ACC-CAA-C-3'

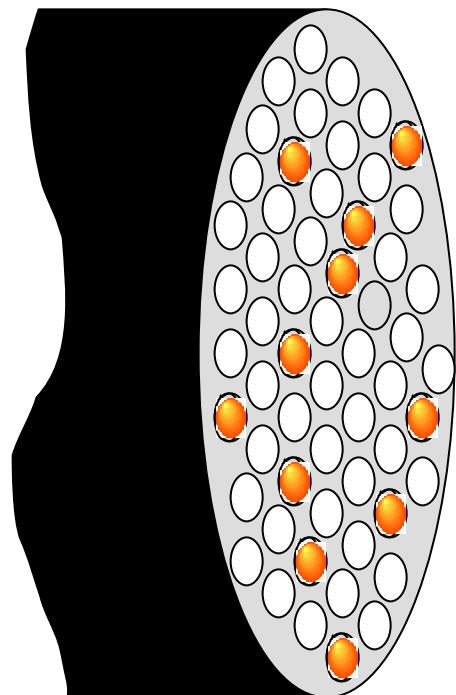
F508C 5'-TA-TCA-TCT-GTG-GTG-TTT-CCT-A-3'

DNA Minimum Hybridization Time with ICCD Camera

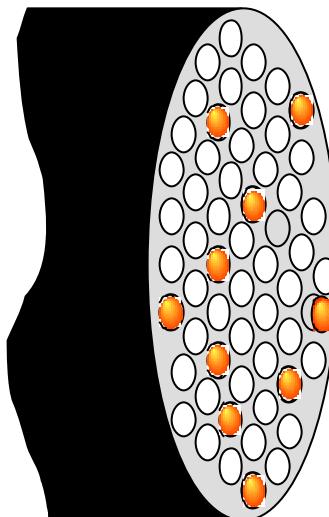
DNA Concentration	Hybridization Time (min)
1 pM	10
100 fM	20
10 fM	30
1 fM	60

Detection Limit Problem

Multiple beads provides a signal averaging benefit.



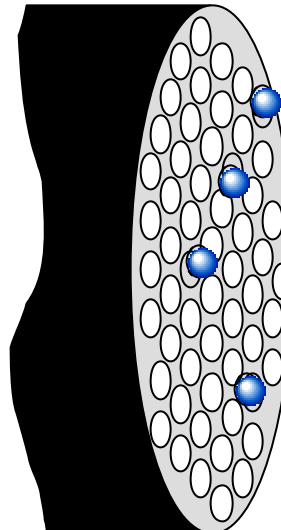
S/N increases by \sqrt{n}



$$\frac{1000 \text{ target molecules}}{10 \text{ beads}}$$

$$=100 \text{ target molecules/bead}$$

Fewer beads provide more target molecule numbers per bead.



$$\frac{1000 \text{ target molecules}}{4 \text{ beads}}$$

$$=250 \text{ target molecules/bead}$$

Multiplexed Array Sensitivity and Selectivity with 1 fM IL2 Target Solutions

IL2 Target - 1 fM concentration - 12 hour hybridization time

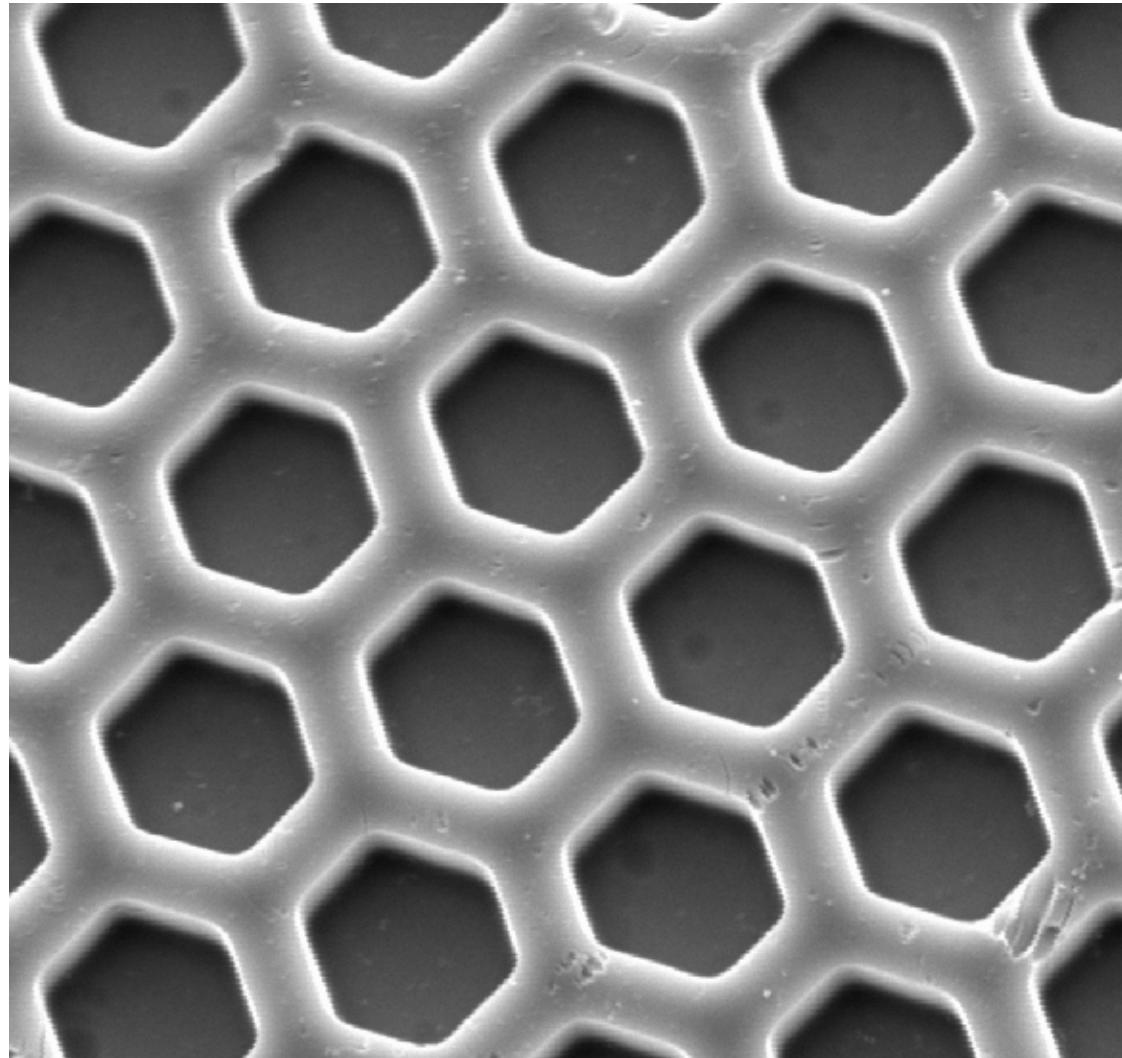
Target/Probe		Mean background \pm s.d	Hybridization \pm s.d.	Signal \pm s.d
F508	F508C	530.43 ± 1.8	550.17 ± 7.5	$\{19.74\} \pm 7.7$
	IL2	563.99 ± 7.7	677.08 ± 8.1	113.09 ± 11
	IL6	445.99 ± 3.9	449.16 ± 1.4	$\{3.17\} \pm 4.1$
IL2	F508C	439.64 ± 3.5	443.34 ± 5.6	$\{3.70\} \pm 6.6$
	IL2	432.52 ± 5.6	503.31 ± 6.6	70.79 ± 8.7
	IL6	431.11 ± 2.1	432.13 ± 2.8	$\{1.02\} \pm 3.5$
IL6	F508C	454.84 ± 3.6	465.82 ± 1.4	$\{10.98\} \pm 3.8$
	IL2	429.42 ± 0.92	517.38 ± 2.6	87.96 ± 2.8
	IL6	459.81 ± 3.0	467.82 ± 5.3	$\{8.01\} \pm 6.1$

Microsphere Array Sensitivity and Selectivity with 100 aM IL2 Target Solutions

IL2 Target - 100 aM concentration - 12 hour hybridization time

Probe/Target		Mean background \pm s.d	Hybridization \pm s.d.	Signal \pm s.d
IL2	F508C	386.97 ± 3.2	387.98 ± 1.4	$\{1.01\} \pm 3.5$
	IL2	378.55 ± 2.3	394.00 ± 3.7	15.32 ± 4.4
	IL6	382.80 ± 6.3	393.81 ± 6.1	$\{11.01\} \pm 7.1$
IL2	F508C	268.66 ± 2.3	274.22 ± 8.5	$\{5.56\} \pm 8.8$
	IL2	297.73 ± 2.3	310.02 ± 2.3	12.29 ± 3.2
	IL6	247.59 ± 2.7	248.70 ± 6.9	$\{1.11\} \pm 7.4$
IL2	F508C	410.73 ± 2.6	413.63 ± 2.6	$\{2.90\} \pm 2.9$
	IL2	410.69 ± 2.7	455.26 ± 6.5	44.57 ± 7.0
	IL6	390.24 ± 7.4	392.88 ± 2.8	$\{2.64\} \pm 7.9$

SEM of a Microwell Array

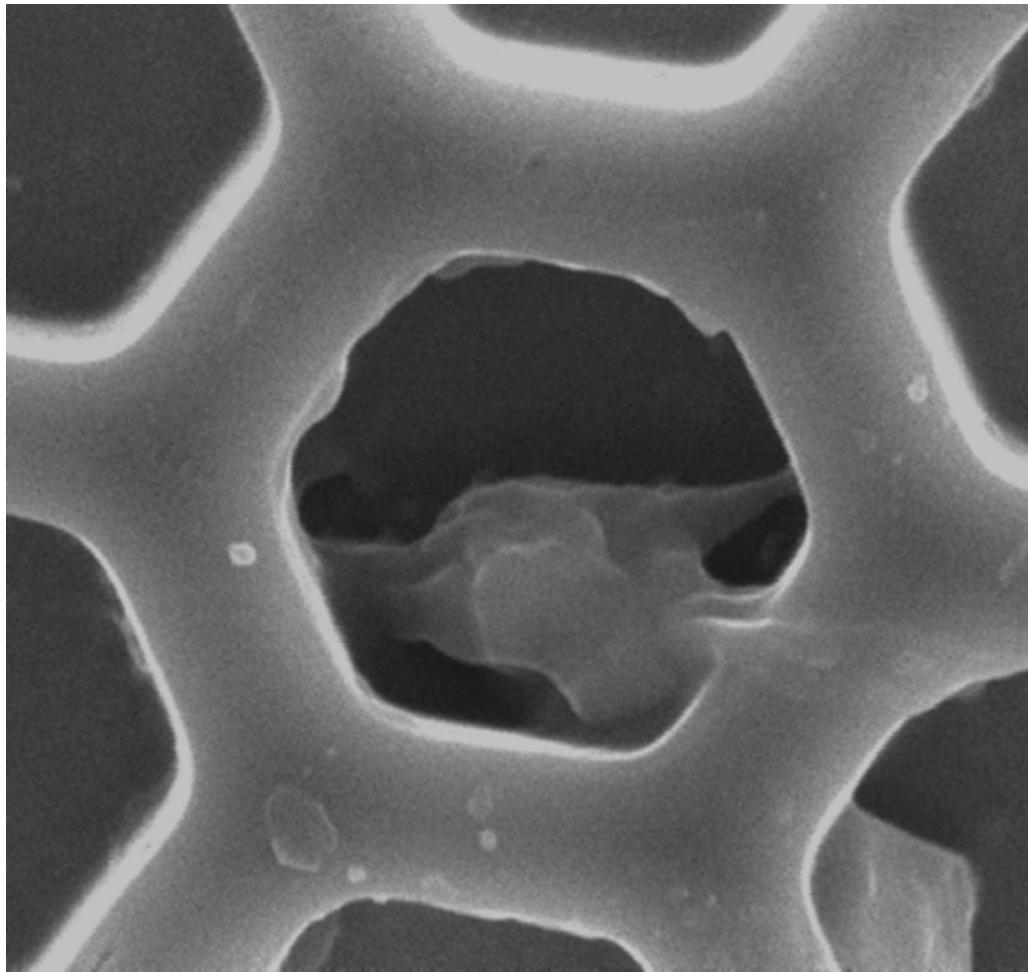


7 μ m well diameter

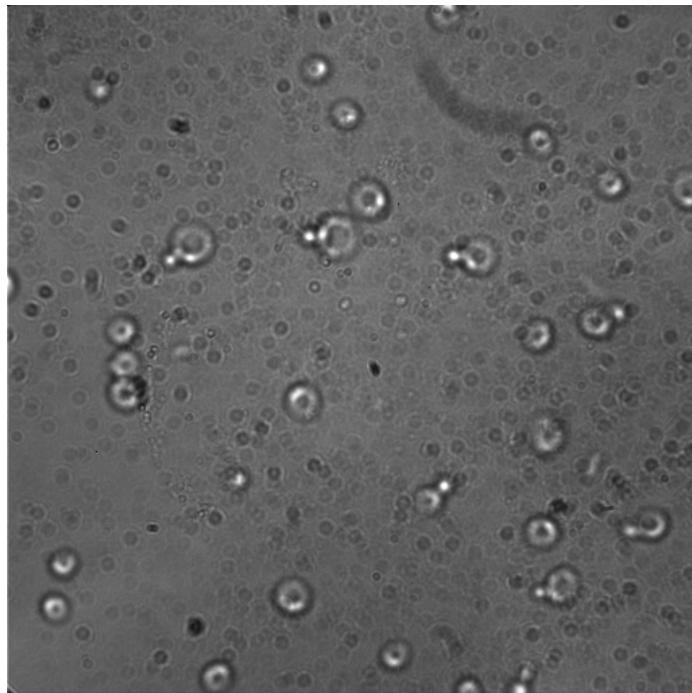
~3 μ m well depth

~90 fL well volume

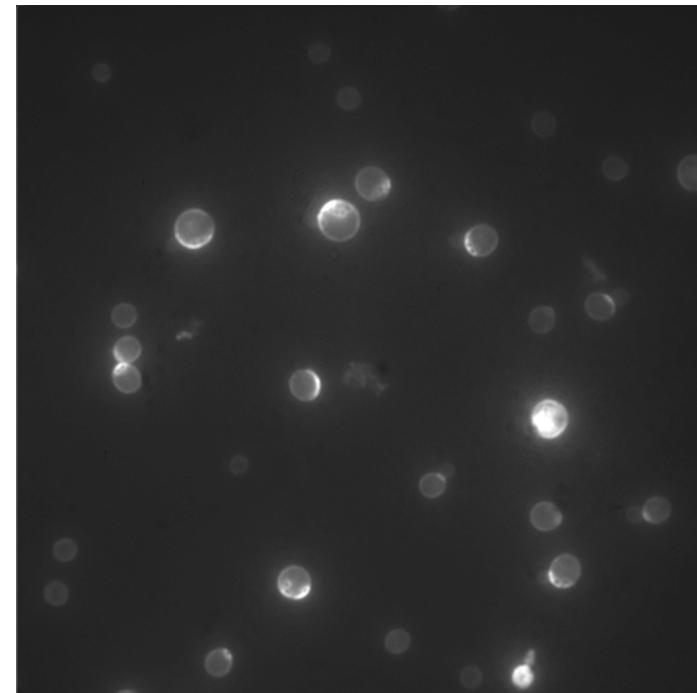
Single NIH 3T3 Mouse Fibroblast Cell in a Fiber-optic Microwell



Single Yeast (*Saccharomyces cerevisiae*) Cells Array

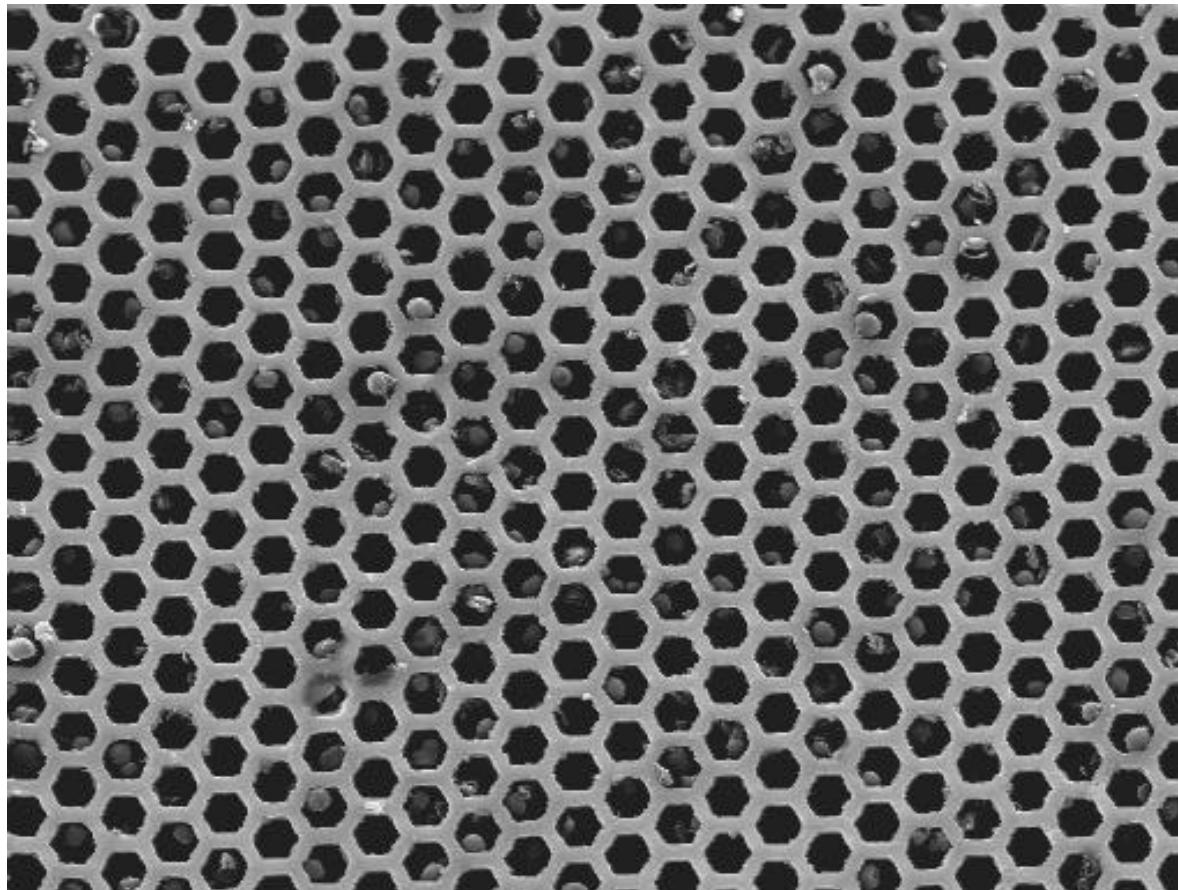


White light

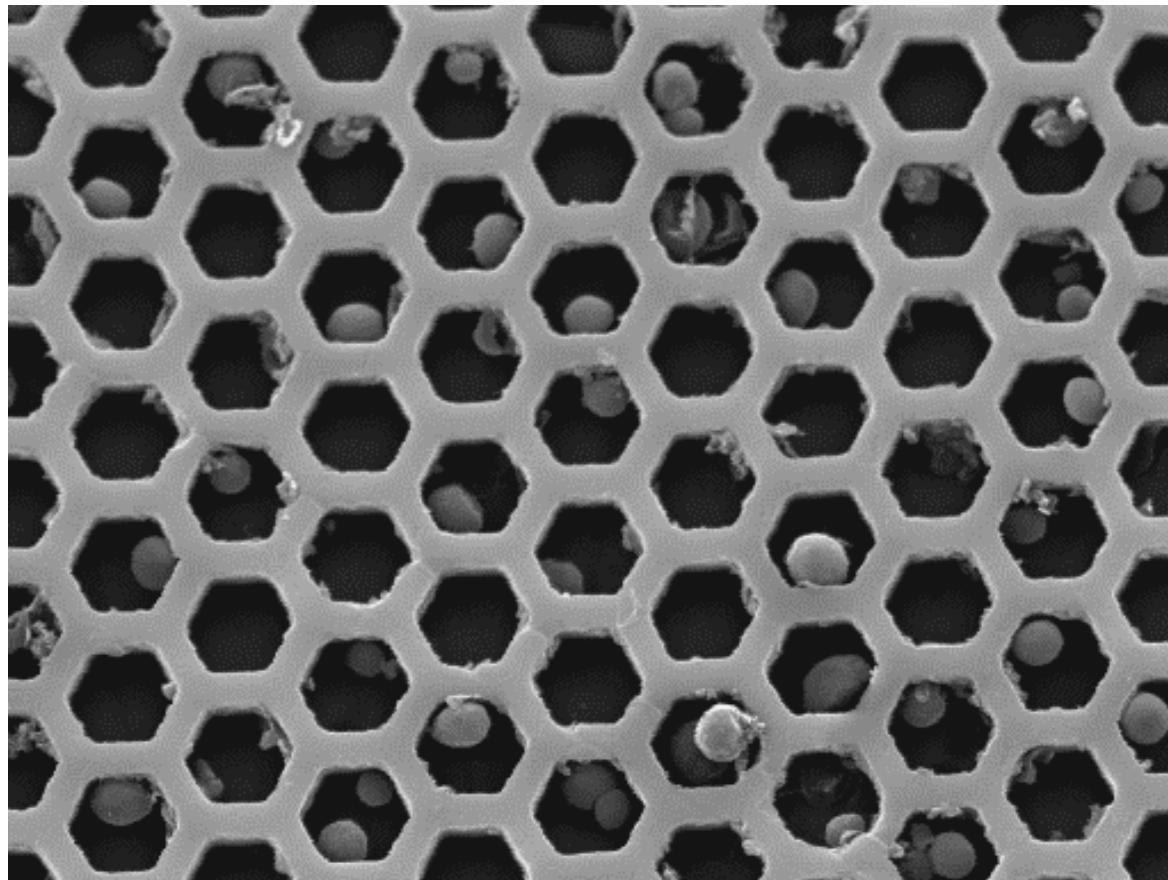


Calcofluor White 360/440

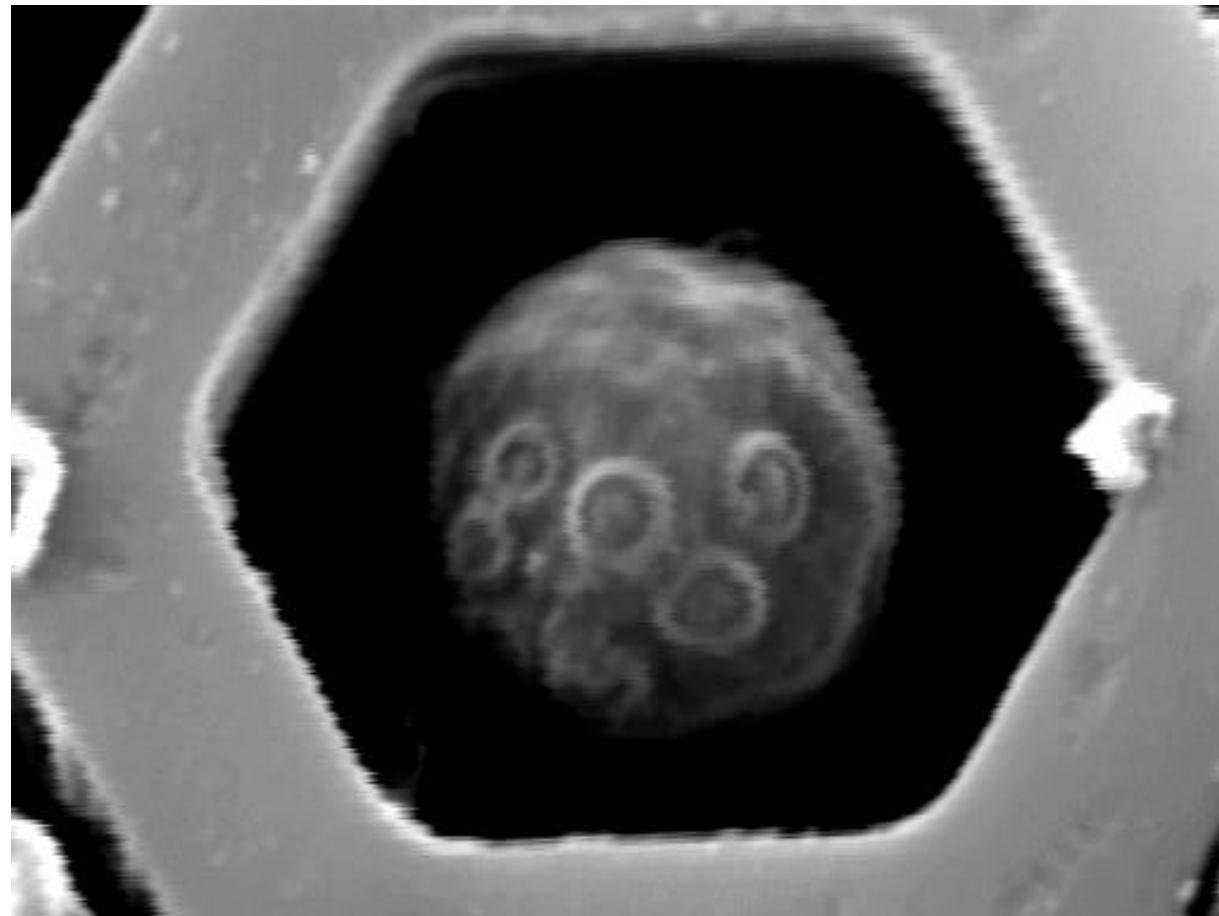
SEM images of Single Yeast Cells on the Microwells array



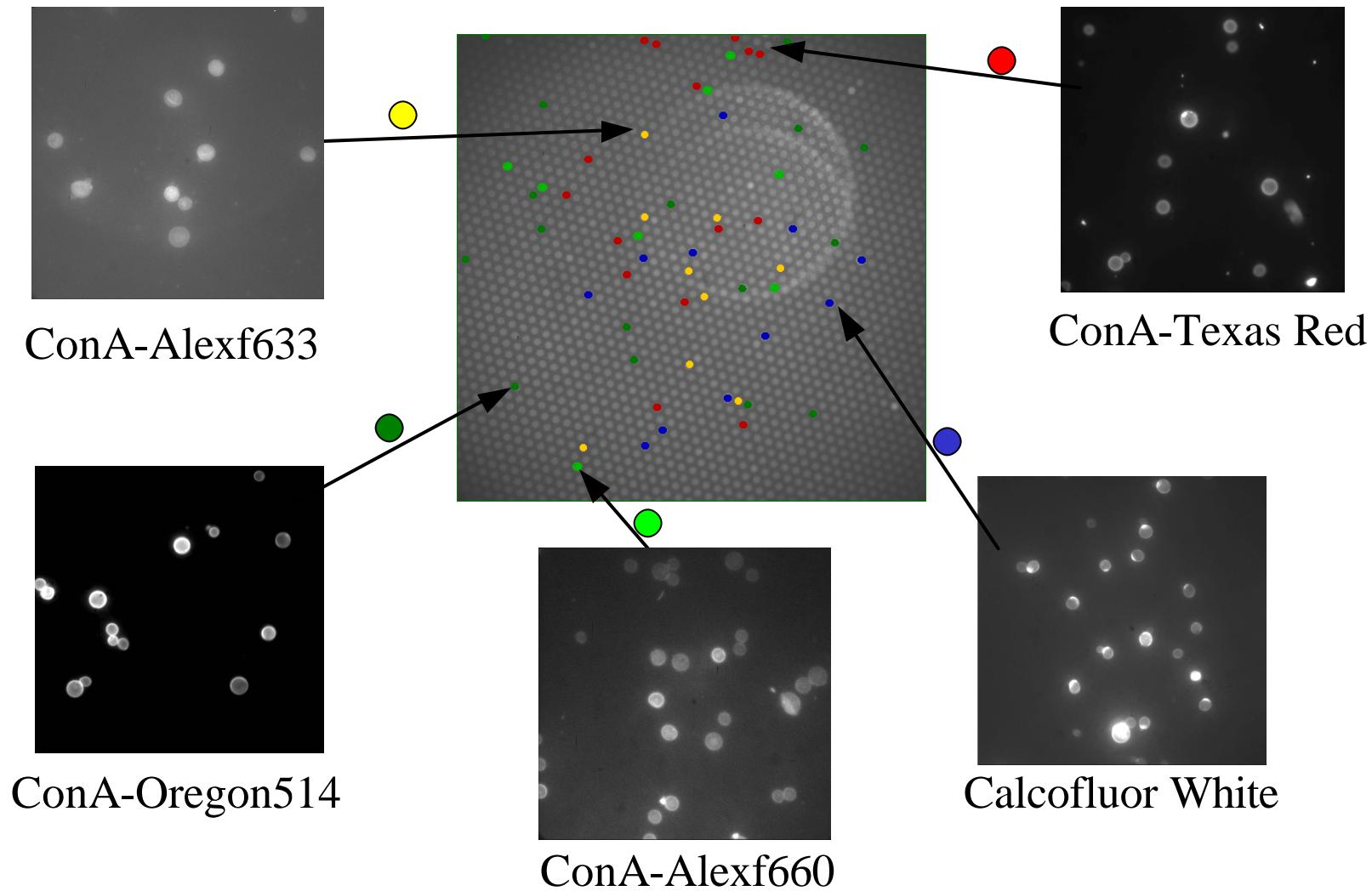
SEM images of Single Yeast Cells on the Microwells array



SEM images of Single Yeast Cells on the Microwells array

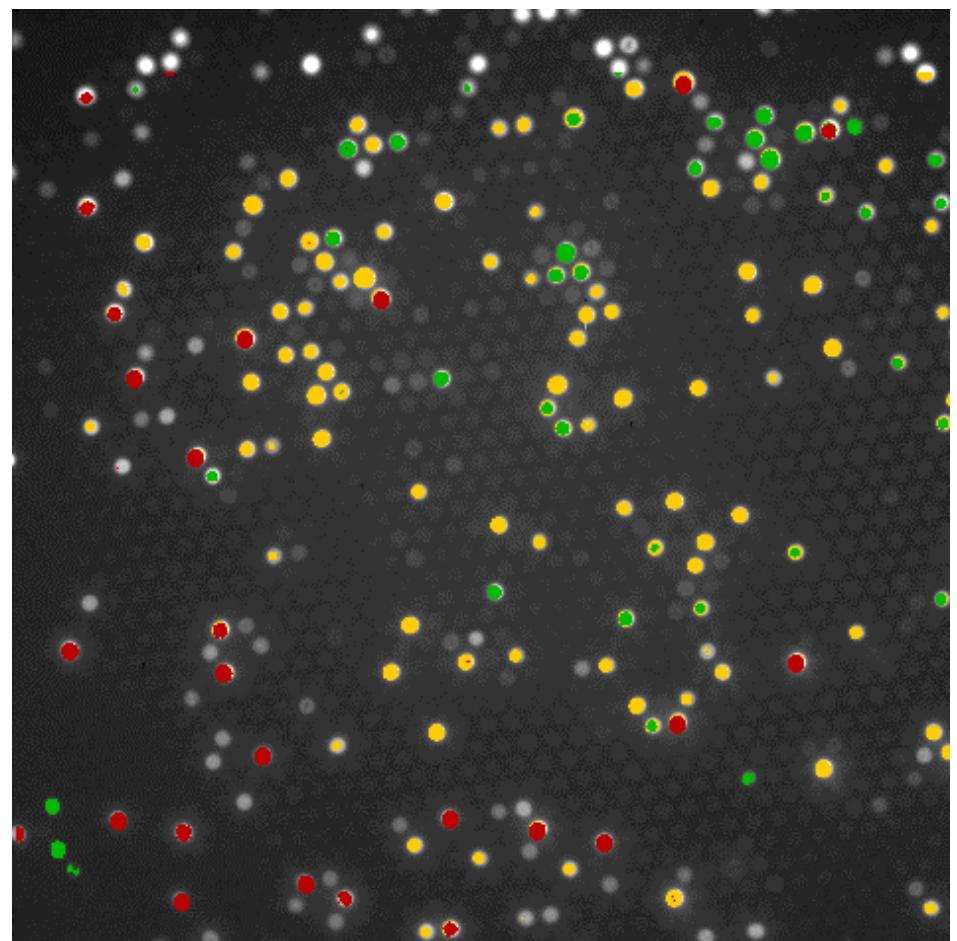


Encoded Yeast Cells on the Fiber Array



pH Measurement of Single Yeast Cells Microenvironment in the Array

- Concanavalin A-FITC
- Concanavalin A-FITC
+
Concanavalin A-Alexa fluor 660
- Concanavalin A-FITC
+
Concanavalin A-Texas Red



Smarter Sensors- Anticipatory

Is it bad?

What does it resemble?

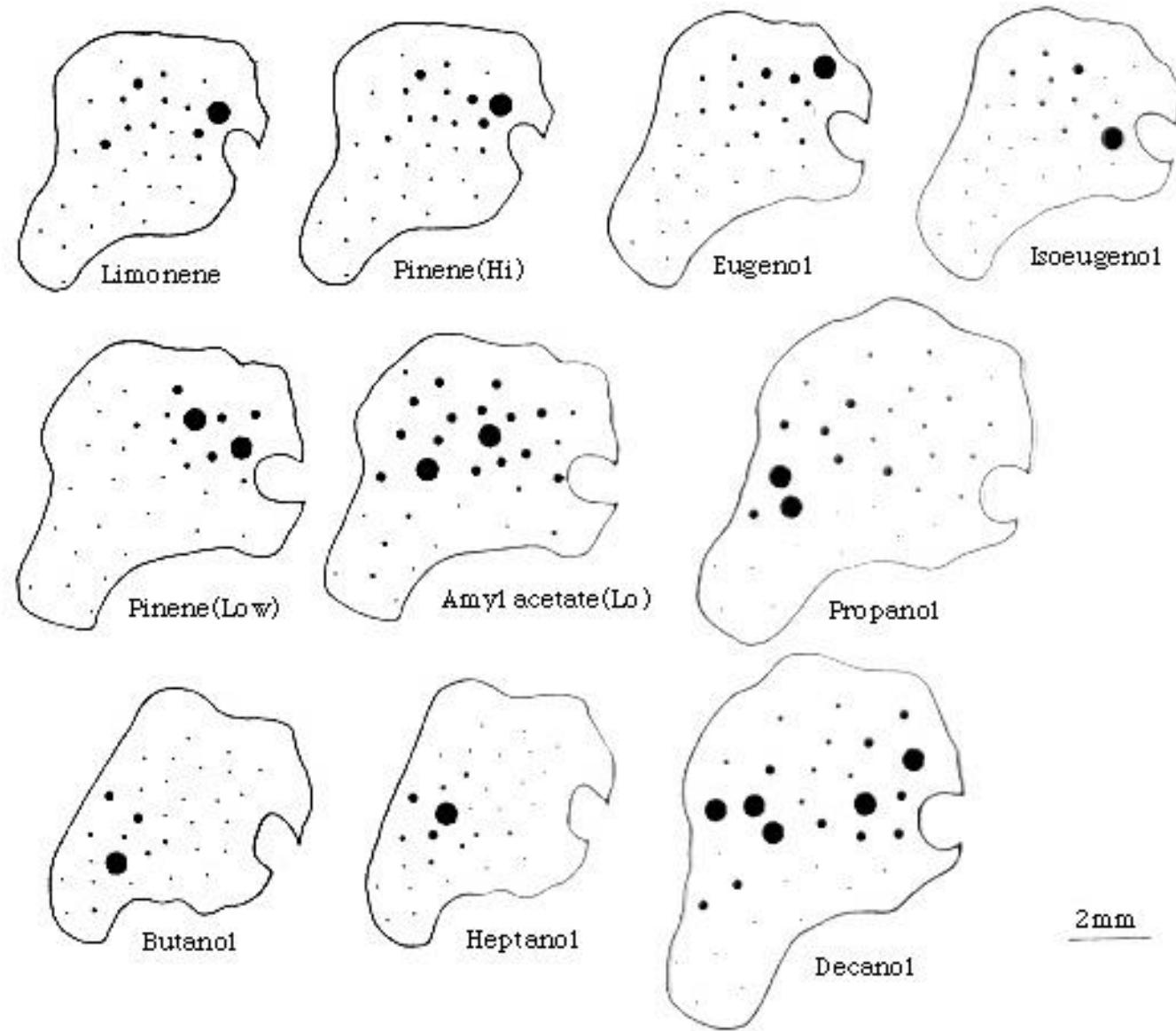
What will it do?

e.g. GI, neurotoxic, etc.

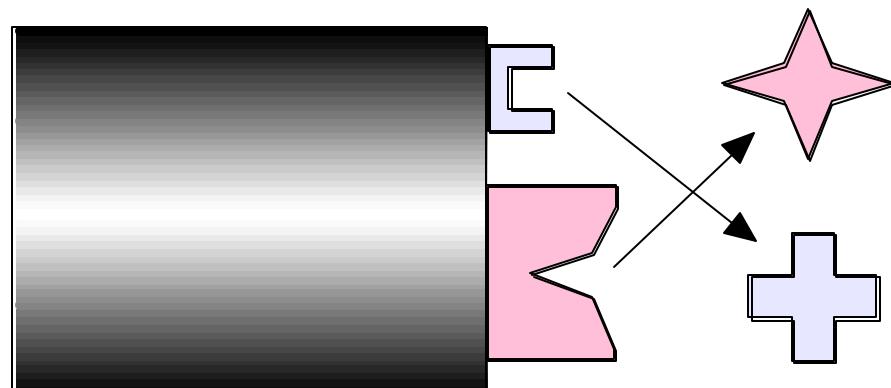
“common virulence mechanisms”



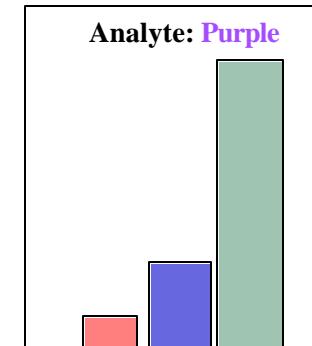
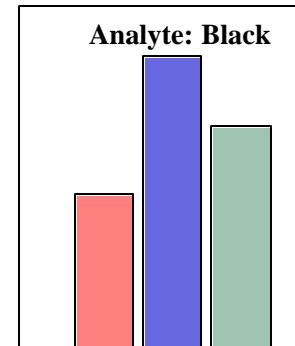
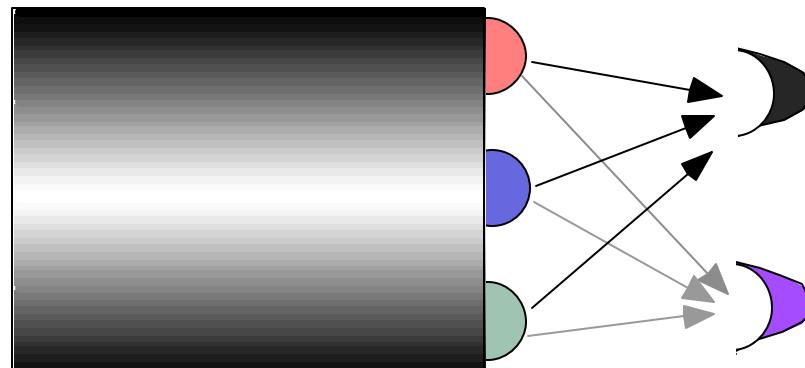
surrogates



Sensor Design

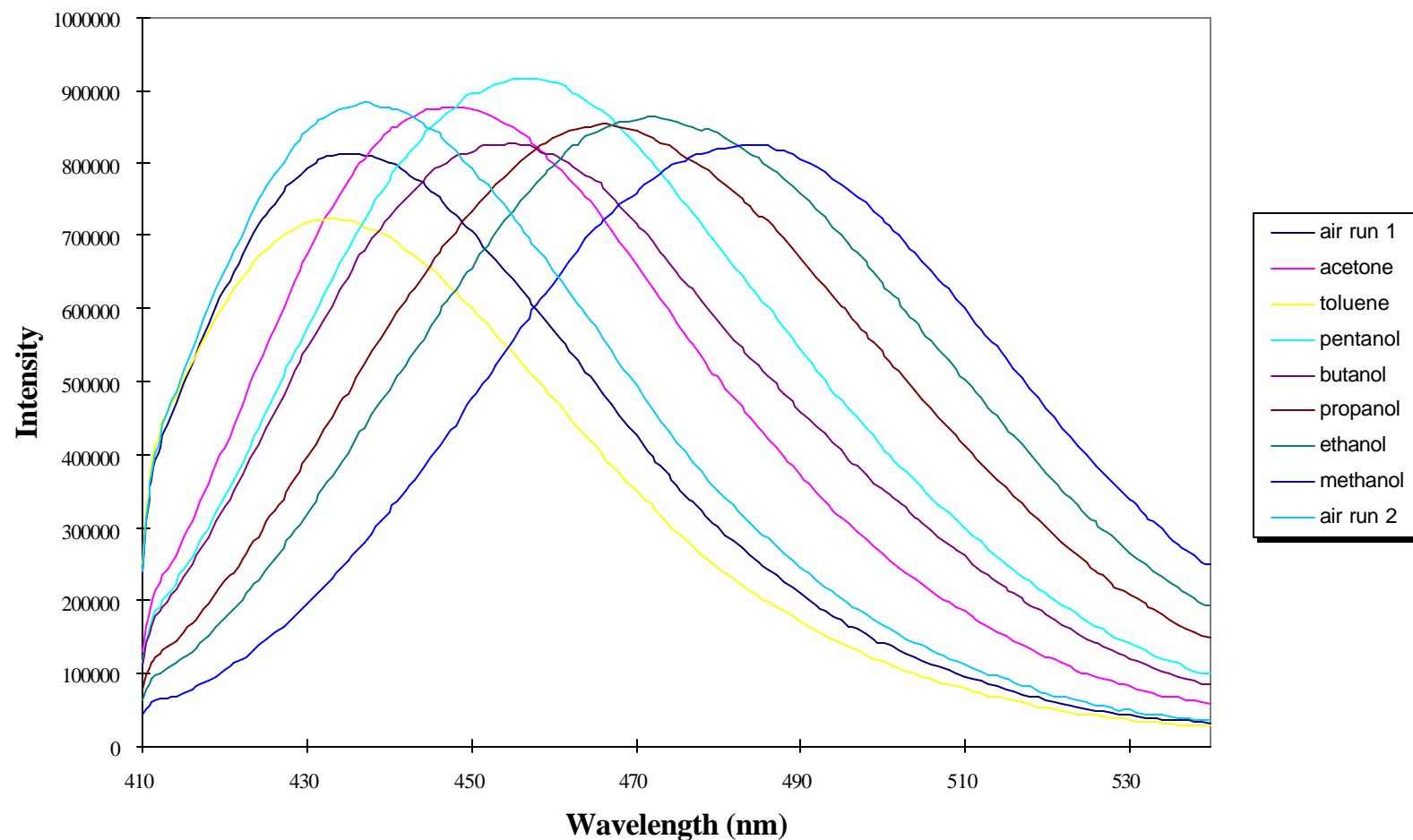


A) Lock-and-key Sensor

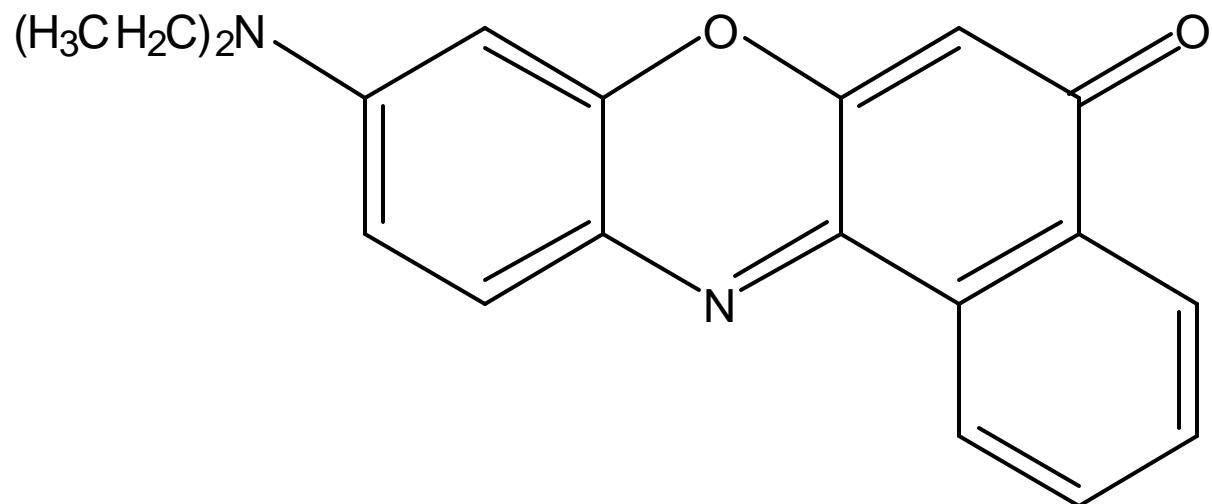


B) Cross-reactive Sensor

Solvatochromic Effect

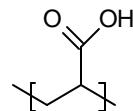


Nile Red

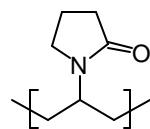


Role of Polymer Polarity

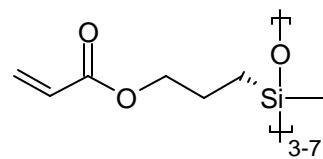
poly(acrylic acid), **PAA**



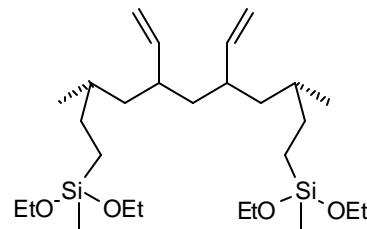
poly(N-vinyl pyrrolidone), **PVP**



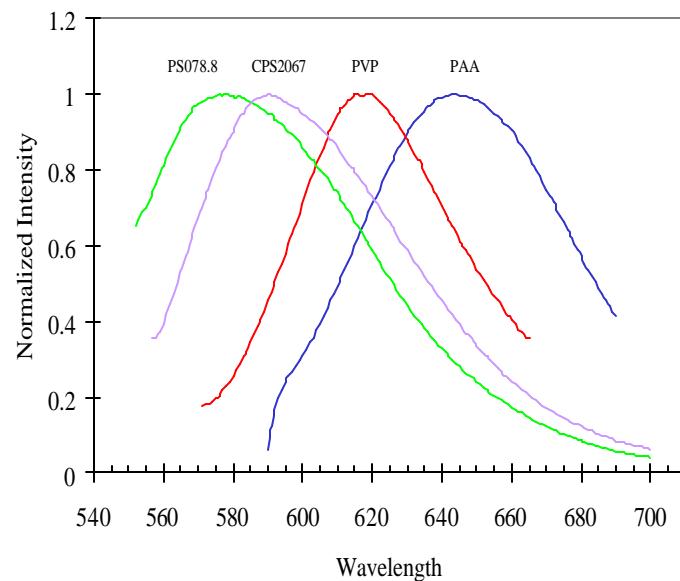
acryloxypropylmethyl-cyclosiloxane, **CPS2067**



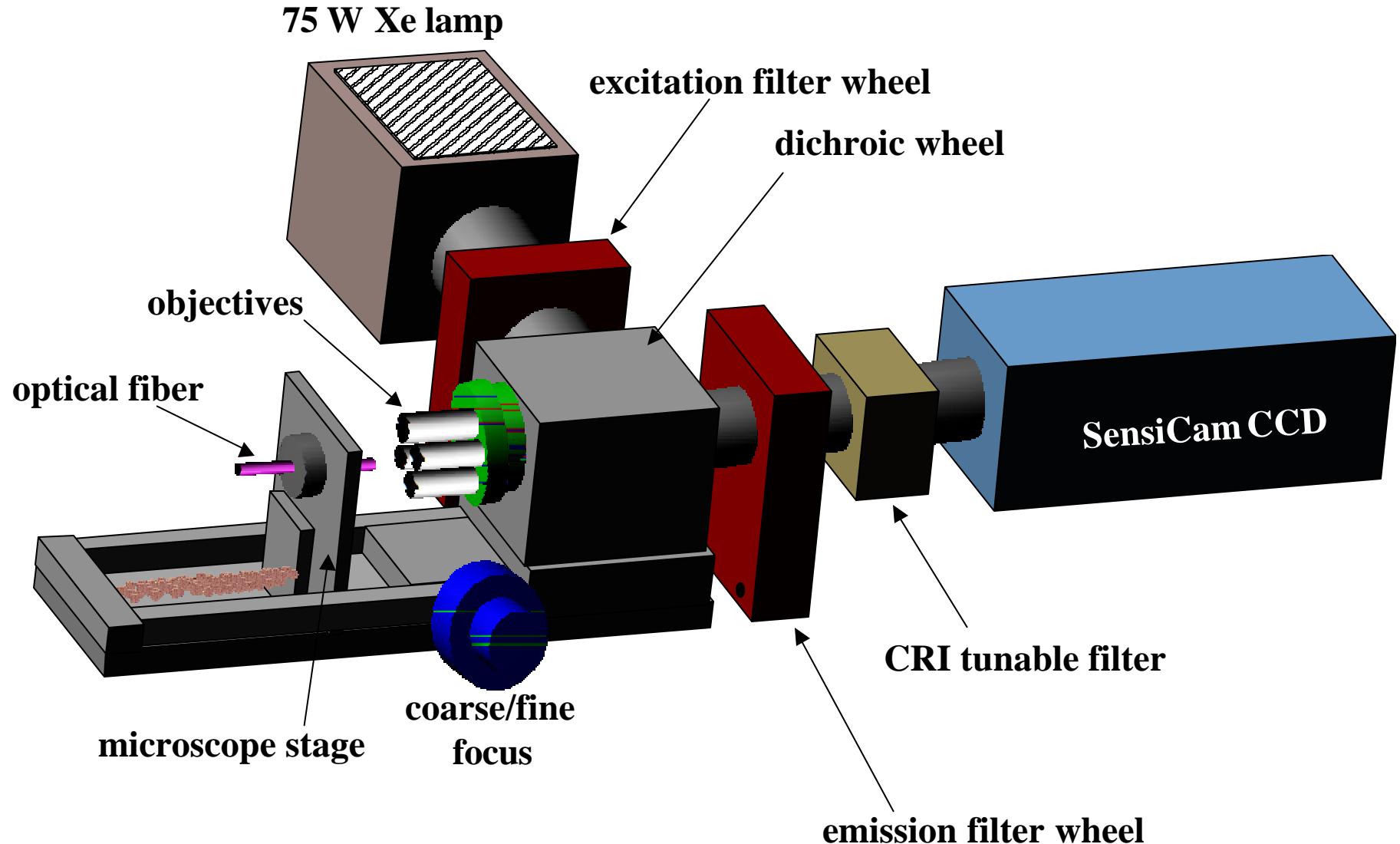
diethoxymethylsilyl-modified polybutadiene, **PS078.8**



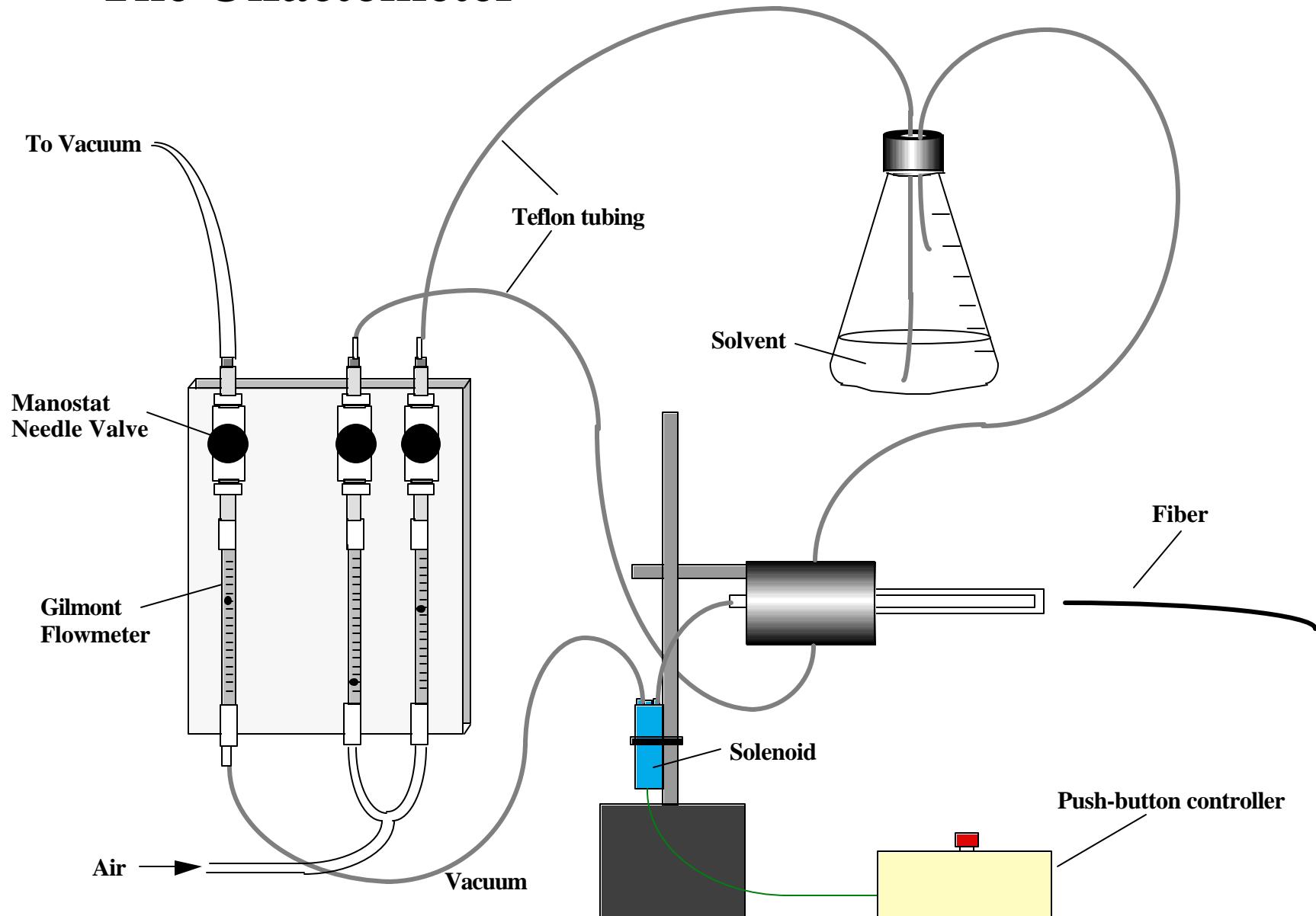
Decreasing polarity



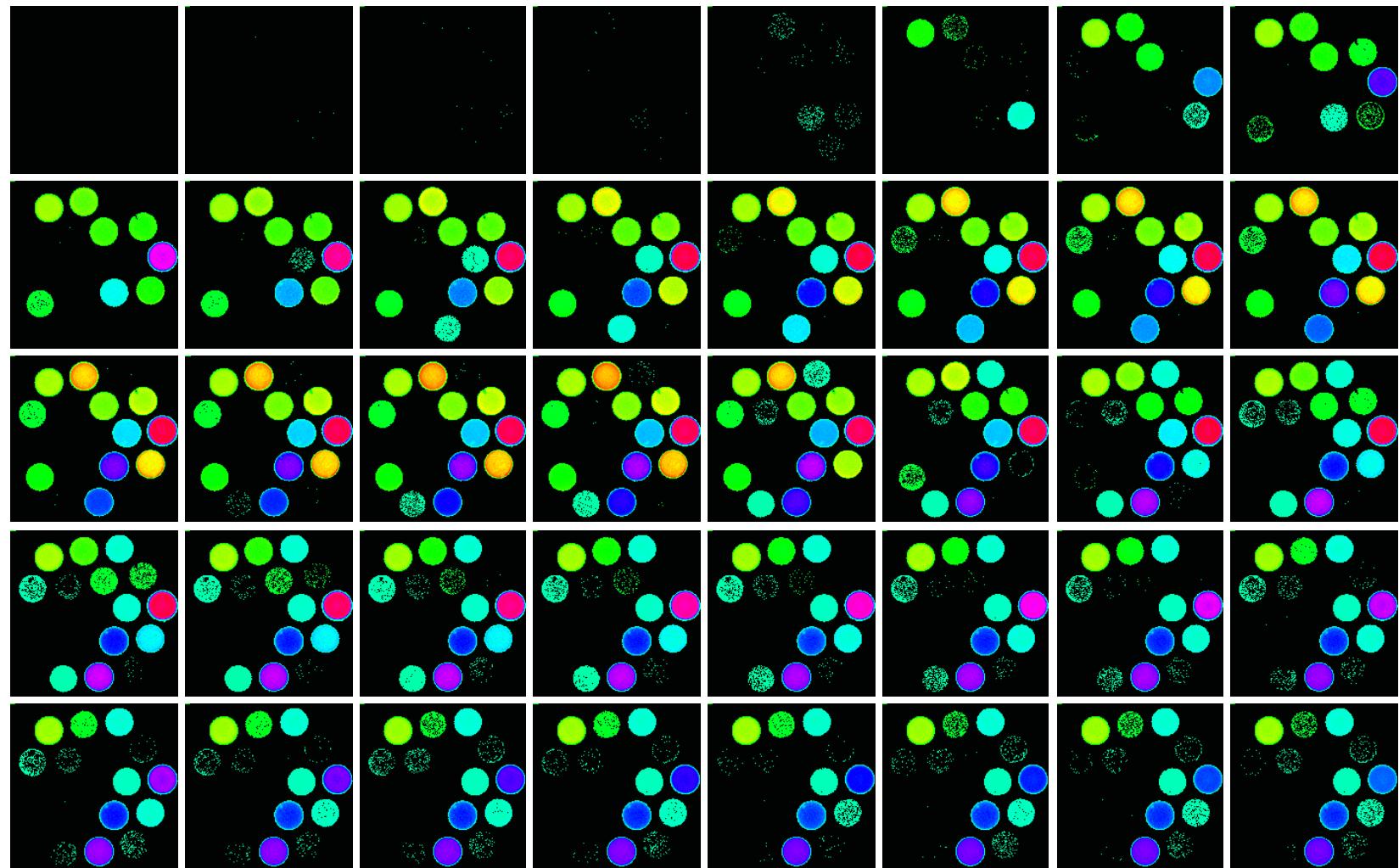
CCD-based imaging system



The Olfactometer — J. Kauer

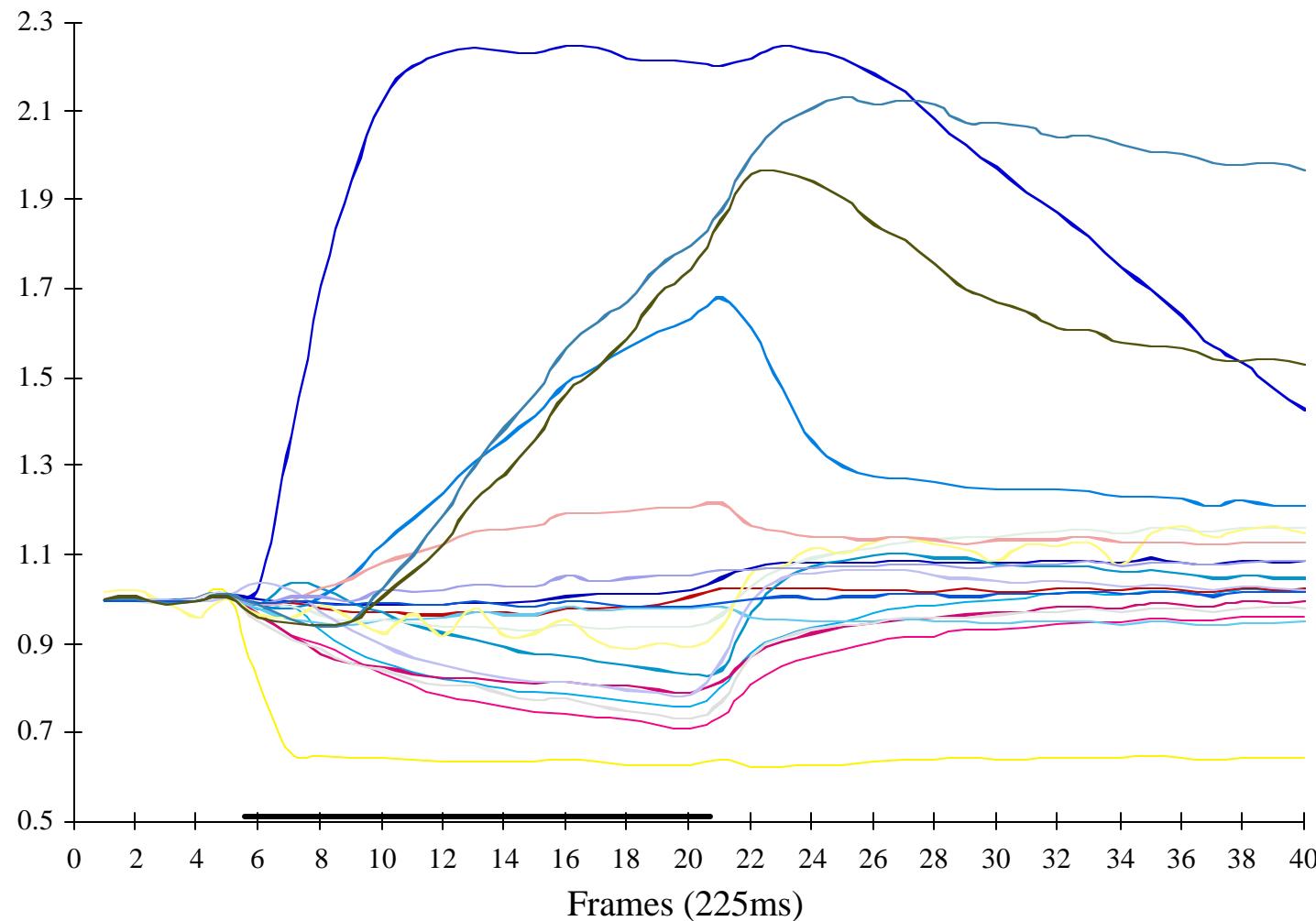


Sensor Array Response to Benzene Vapor Pulse



Dickinson, T. A., et al. (1996) *Nature*, 382: 697-700.

Temporal Plots from 19-Fiber Sensor Array Response to Benzene Vapor Pulse



Classification Results

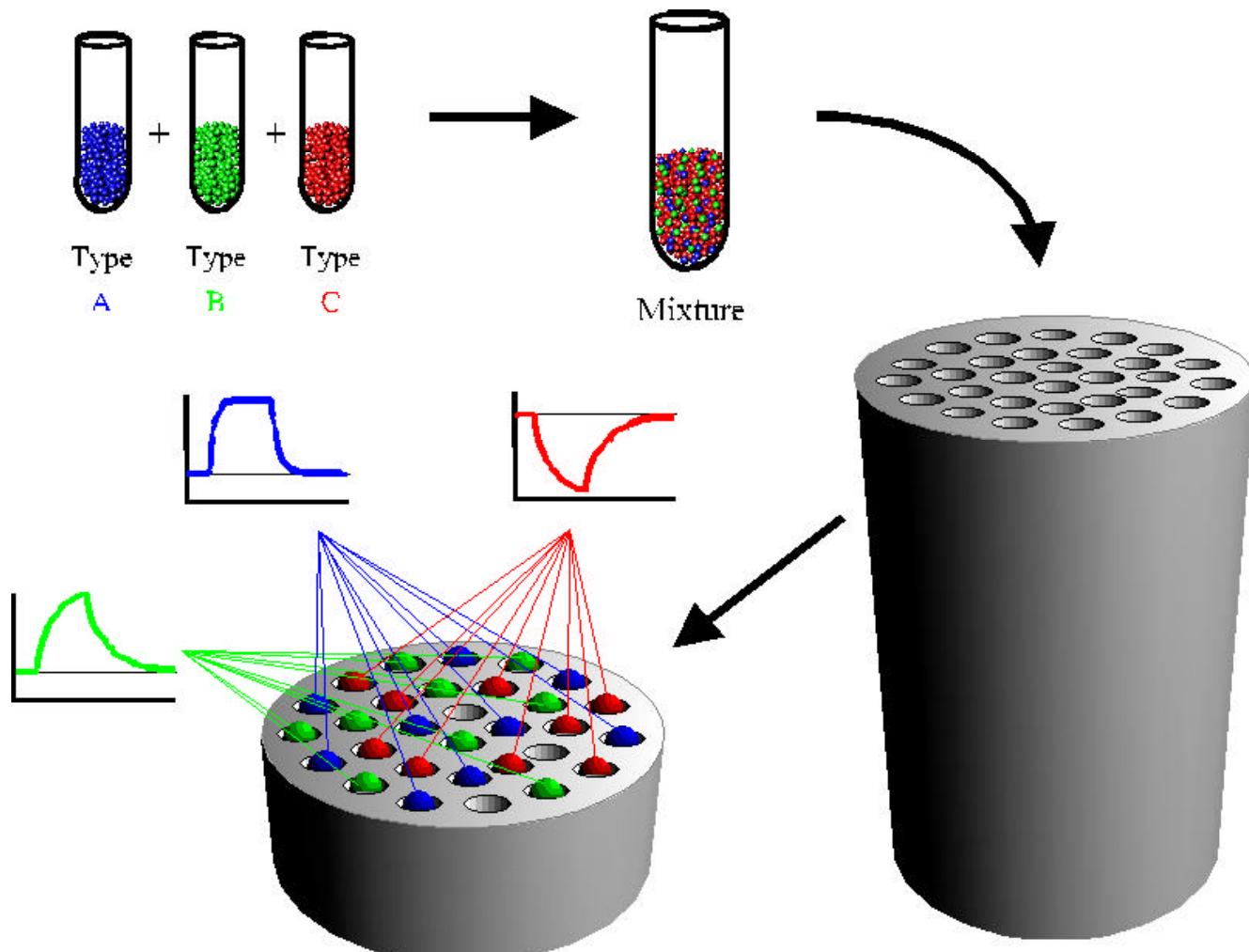
Learning Vector Quantization Approach

True
Identity

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20
1	24																			
2		23																		
3			20								3			1						
4				1	19						1								2	
5				1		17	4				2									
6					1	2	19					1						1		
7							18	4			1			1						
8							2	21						1						
9					1				23											
10					3					19			1				1			
11										21						1		2		
12											23							1		
13									1			23								
14									1				20			2		1		
15				1									23							
16													24							
17														24						
18														24						
19									4						20					
20		1																23		

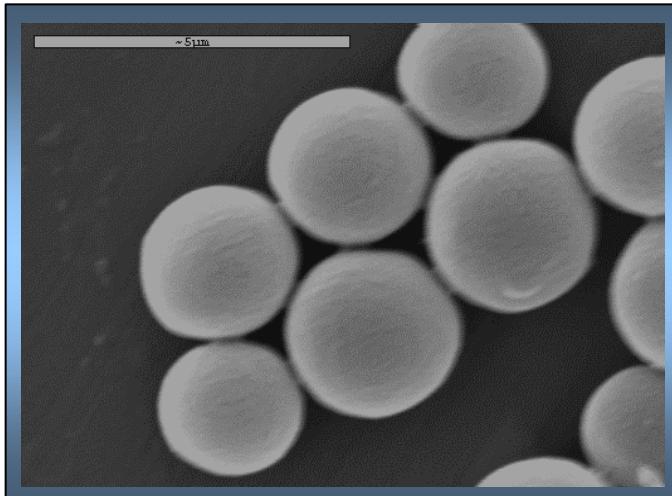
- 1) Acetone
- 2) Butyl Acetate
- 3) Beauty
- 4) Camphor
- 5) Carvone -
- 6) Carvone +
- 7) Chloroform
- 8) Dichloroethane
- 9) DMSO
- 10) Drakkar Noir
- 11) Water
- 12) Heptane
- 13) Isopropanol
- 14) Indole
- 15) Mercaptoethanol
- 16) Methanol
- 17) Propanol
- 18) Propionic Acid
- 19) Pseudoexplosive
- 20) Toluene

SENSOR ARRAYS are Assembled 'Randomly' in ONE Fabrication Step

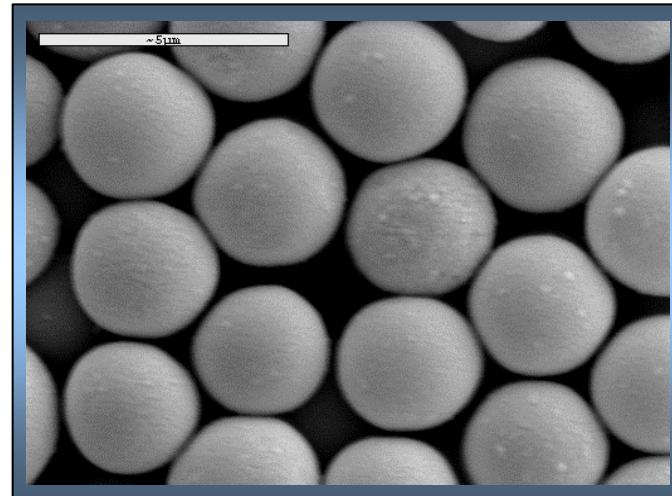


NOTE: the Sensor Array is a 'Self-Encoding' Bead Array (SEBA). Billions of Sensors are Fabricated at Once.

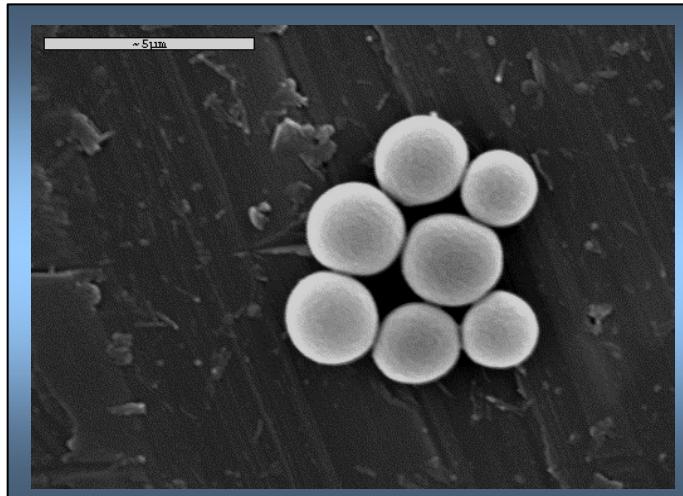
Hollow Poly(benzyl methacrylate) Spheres



3.5 h polymerization



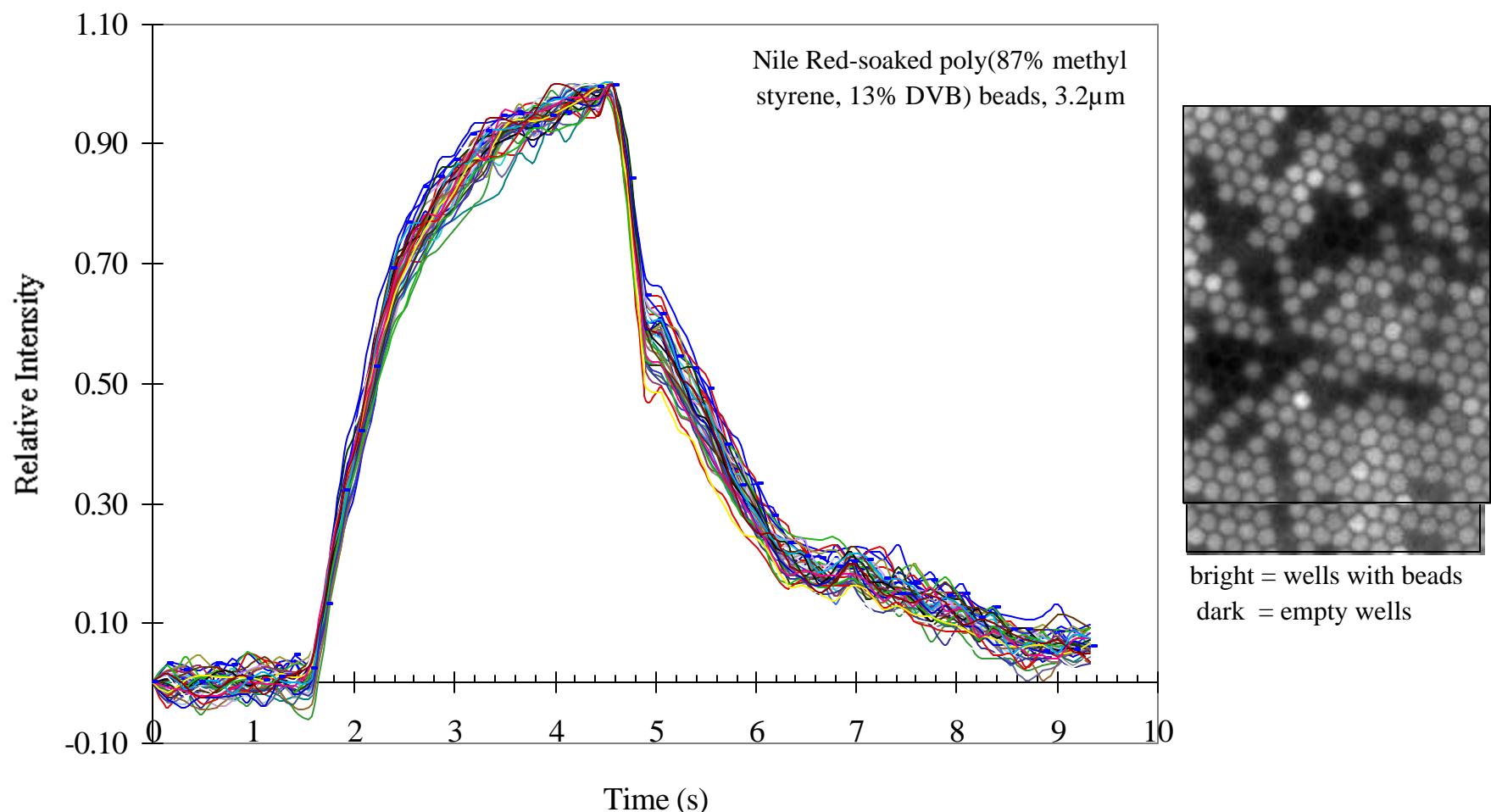
6.5 h polymerization



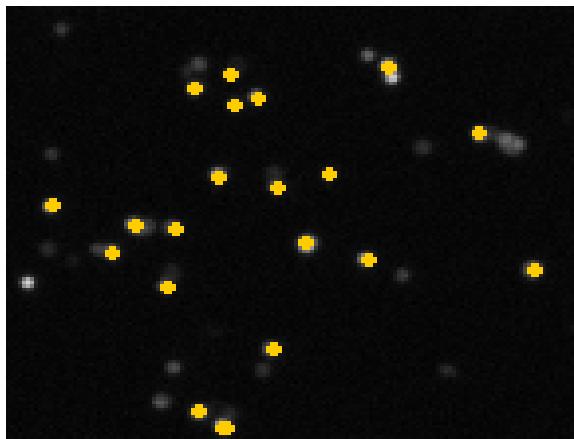
14h polymerization

Chem. Mater. 2000

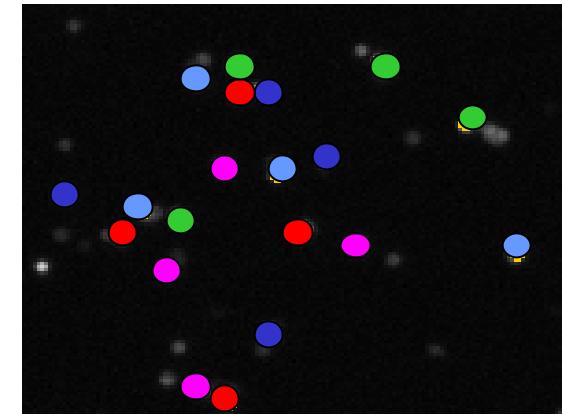
Nile Red/PolyMethylStyrene Beads in Wells: Response of 40 beads to methanol pulse



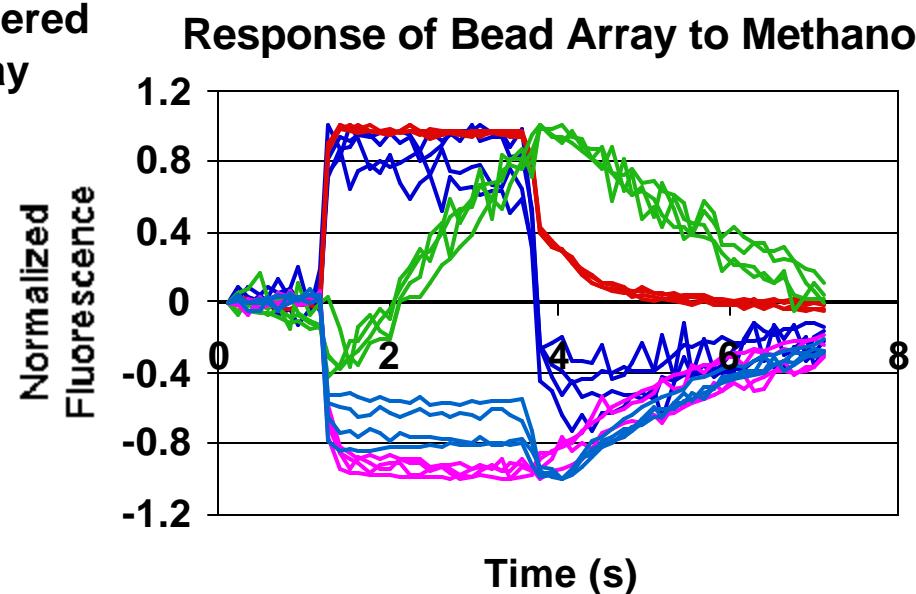
Sensor Registration Problem



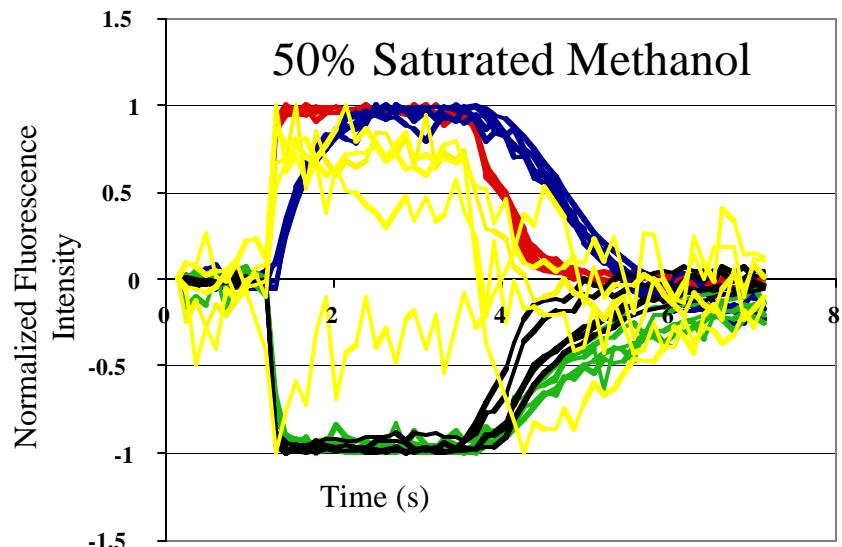
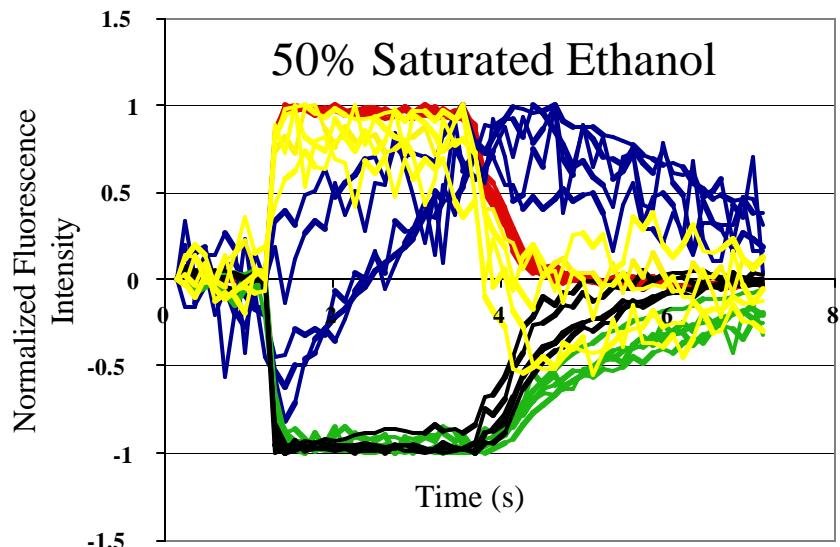
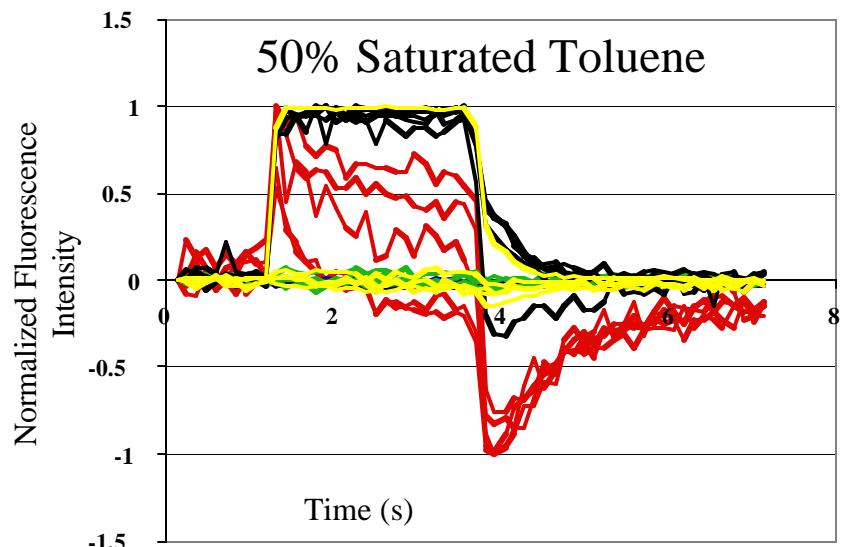
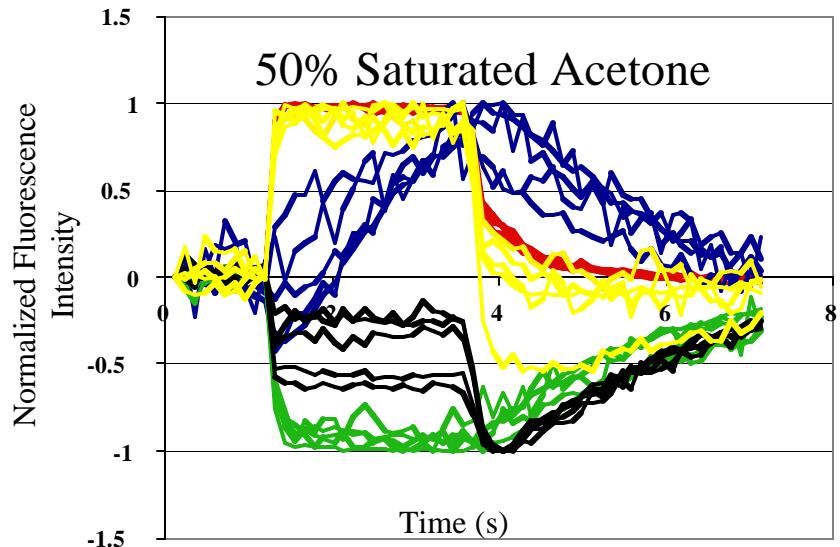
Randomly Ordered
5 Bead Array



Decoded 5
Bead Array

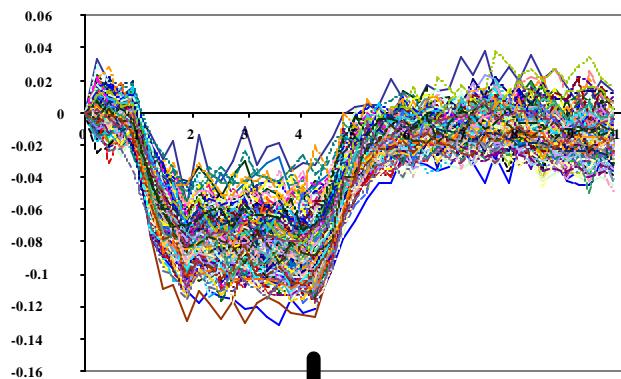


5 Sensor Types with 4 Analytes

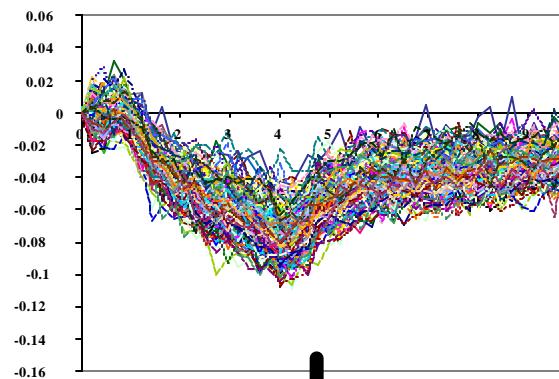


TIME (s) vs. FLUORESCENCE RESPONSE: 250 INDIVIDUAL Bead Sensors

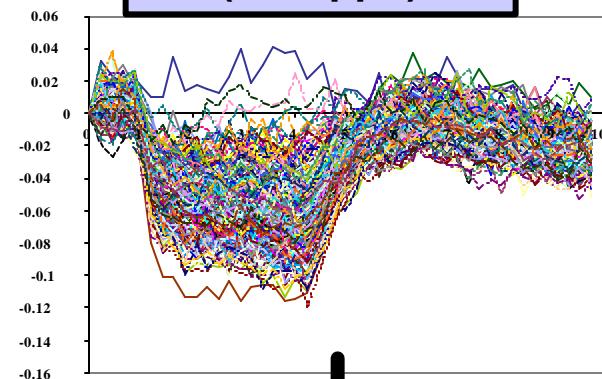
~23 ppb 2,4-DNT



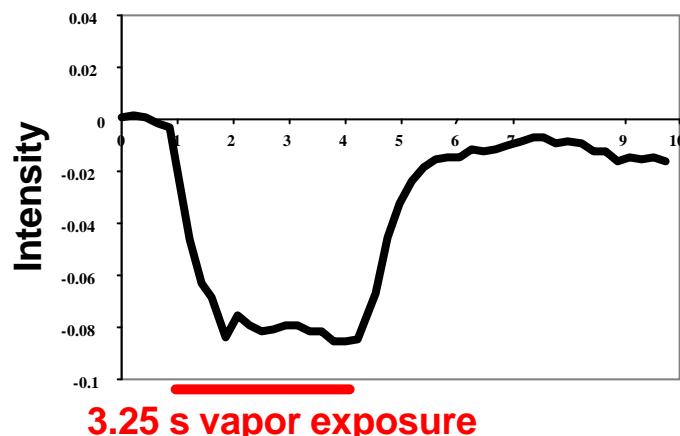
~80 ppb 1,3-DNB



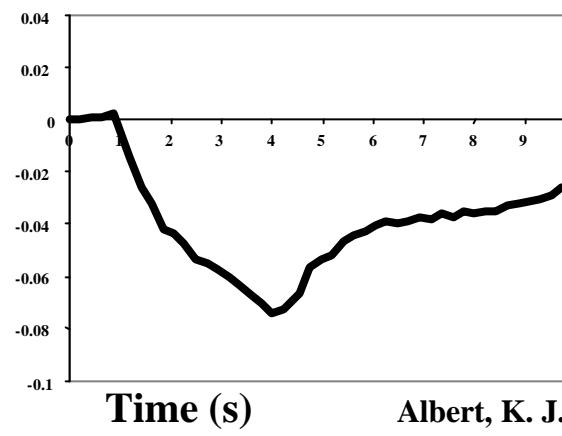
8% saturated
TNT vapor Strips
(~0.4 ppb)



Signal/Noise Improvement:
Average of 1000 Sensors



3.25 s vapor exposure

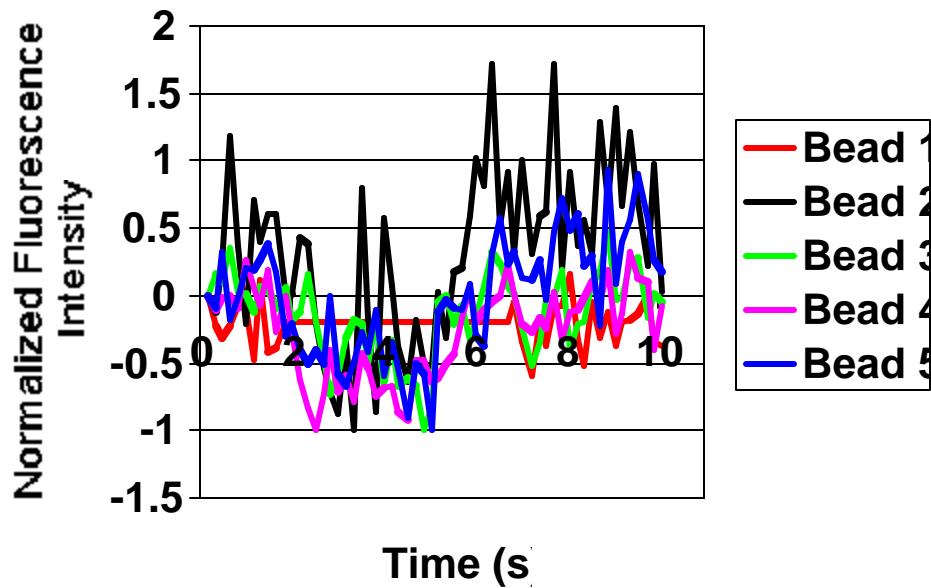


Time (s)

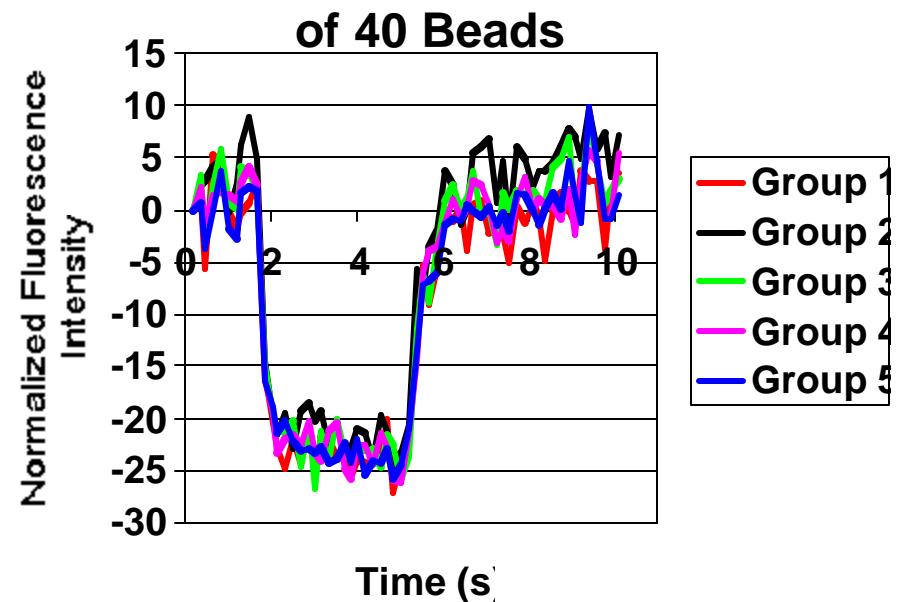
Albert, K. J. and D. R. Walt (2000) *Anal. Chem.* in press.

Signal Summing

Individual Response of 5 Bead Sensors



Summed Responses of 5 Random Groups of 40 Beads



Summing improves signal-to-noise ratio.

Analytes for Two Class Problem

- Pure Analytes
 - Acetone
 - Benzene
 - Chloroform
 - Ethanol
 - Ethyl Acetate
 - Heptane
 - Methanol
 - Toluene
 - 1,3-Dinitrobenzene
 - 4-Nitrotoluene
- Binary Mixtures
 - Ethyl Acetate/Heptane
 - Methanol/Benzene
 - 4-NT/Benzene
 - 4-NT/Heptane
 - 4-NT/Methanol
 - 1,3-DNB/Ethyl Acetate
 - 1,3-DNB/Heptane

Concentrations of Analytes

Table 1: The concentration of the pure analytes $\pm 15\%$.

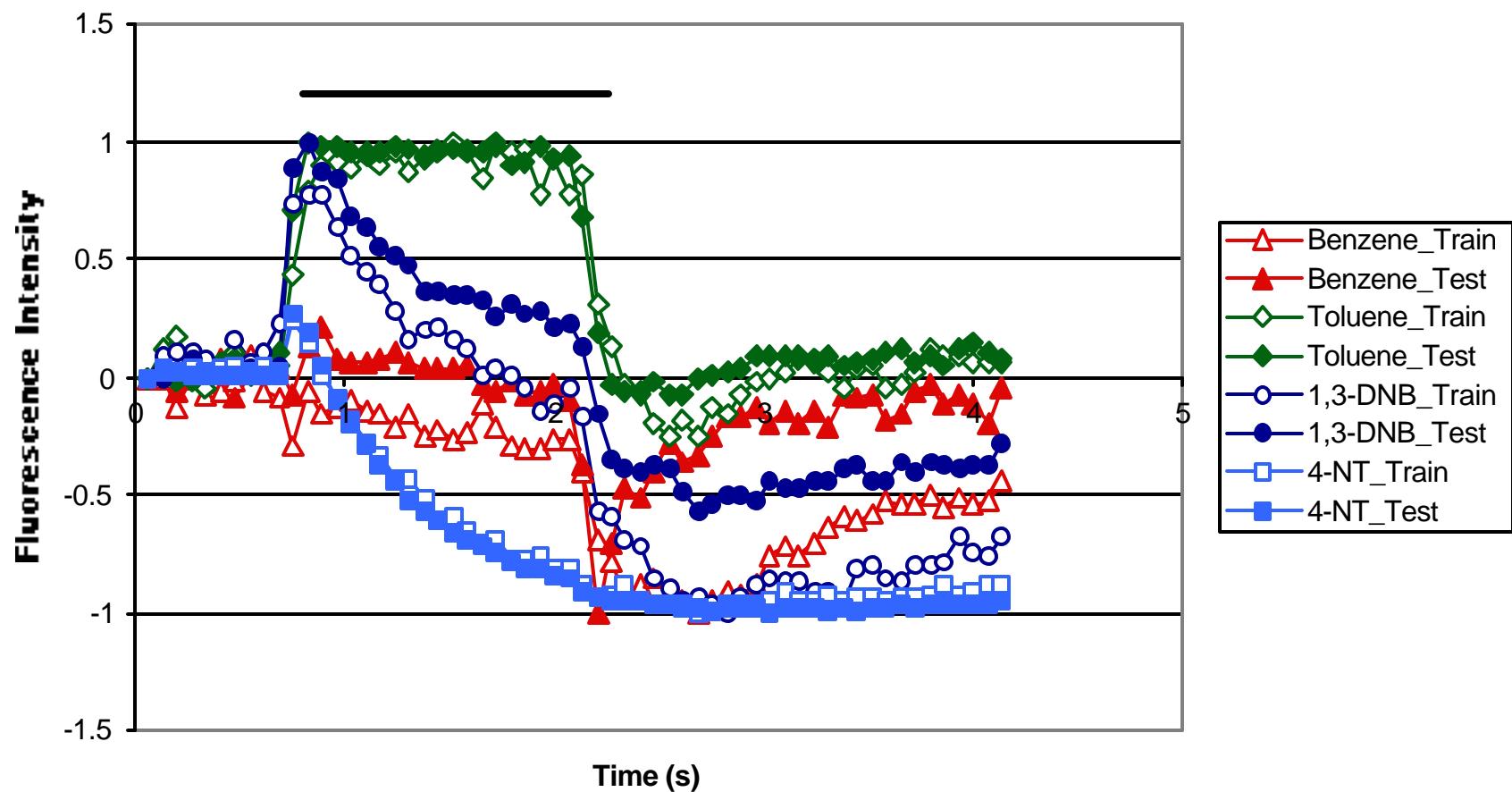
The concentrations were calculated based on literature values for analyte vapor pressures.

Analyte	Vapor Pressure @25 °C (mmHg)	Concentration (ppm)
Acetone	2.31E+02	7.6E+04
Benzene	9.53E+01	3.1E+04
Chloroform	1.97E+02	6.5E+04
Ethanol	5.90E+01	1.9E+04
Ethyl Acetate	9.45E+01	3.1E+04
Heptane	4.57E+01	1.5E+04
Methanol	1.27E+02	4.2E+04
Toluene	2.84E+01	9.4E+03
1,3-Dinitrobenzene	9.00E-04	6.0E-01
4-Nitrotoluene	1.64E-01	1.1E+02

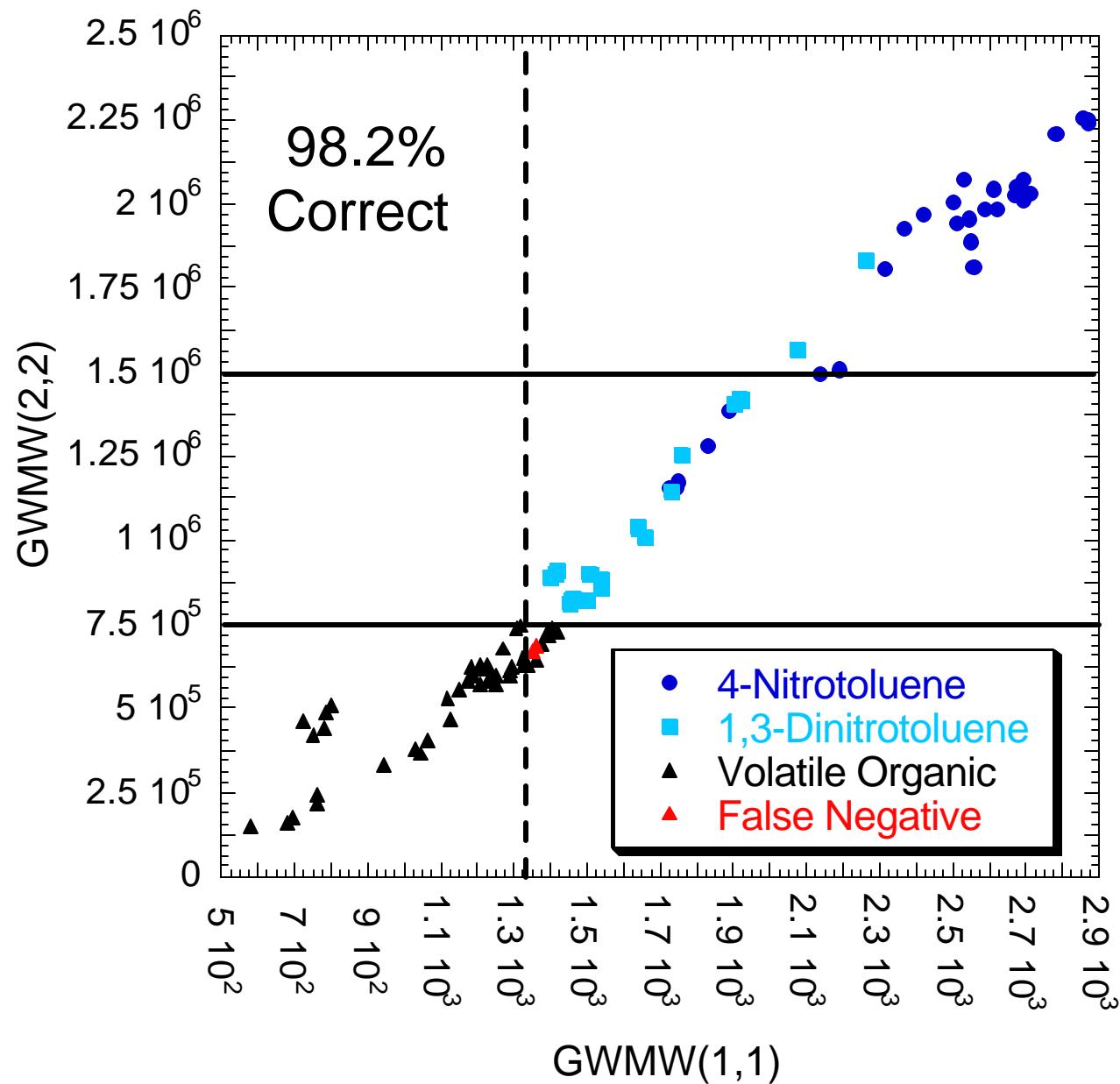
Table 2: The concentration of the binary mixtures $\pm 15\%$.

Analyte 1	Analyte 2	Concentration analyte1 (ppm)	Concentration analyte2 (ppm)
Benzene	Methanol	3.1E+04	4.2E+04
Benzene	4-Nitrotoluene	3.1E+04	5.5E+01
Benzene	4-Nitrotoluene	3.1E+04	1.1E+02
Ethyl Acetate	Heptane	3.1E+04	1.5E+04
Ethyl Acetate	1,3-Dinitrotoluene	3.1E+04	3.0E-01
Ethyl Acetate	1,3-Dinitrotoluene	3.1E+04	6.0E-01
Heptane	1,3-Dinitrotoluene	1.5E+04	6.0E-01
Heptane	4-Nitrotoluene	1.5E+04	1.1E+02
Methanol	4-Nitrotoluene	4.2E+04	5.5E+01
Methanol	4-Nitrotoluene	4.2E+04	1.1E+02

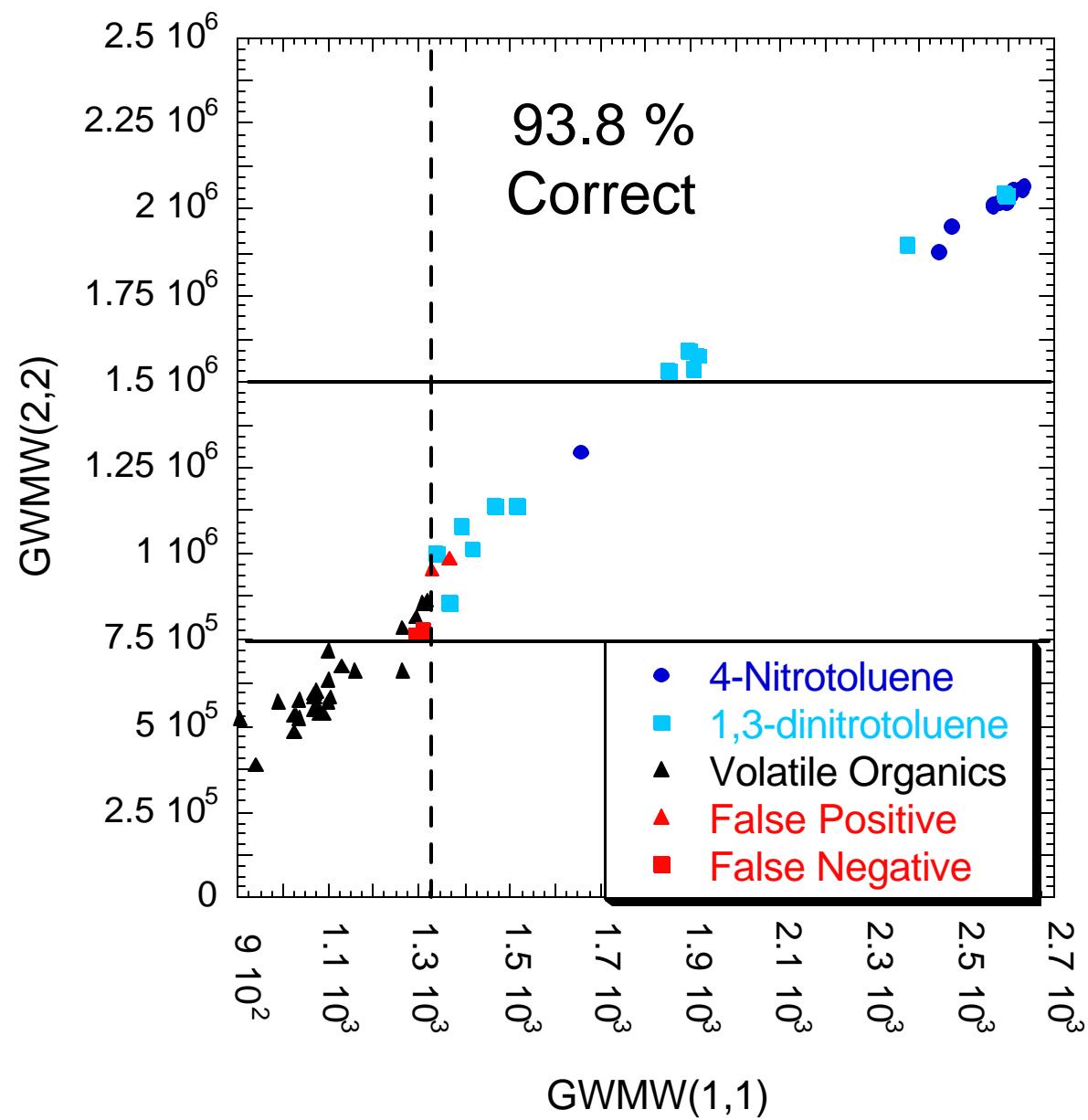
Reproducible Responses from Training to Testing array



First Testing Array (1 Month)



Second Test Array (7 months)



Live/Dead Bacteria Discrimination

Calculated Identity

Actual Identity	Live B10	Live B4	Live B5	Live B8	Live B9	Dead B10	Dead B4	Dead B5	Dead B8	Dead B9	Medium
Live B10	4	1	0	0	0	0	0	0	0	0	0
Live B4	2	3	0	0	0	0	0	0	0	0	0
Live B5	0	0	5	0	0	0	0	0	0	0	0
Live B8	0	0	0	4	0	0	1	0	0	0	0
Live B9	0	0	0	0	5	0	0	0	0	0	0
Dead B10	0	0	0	0	0	5	0	0	0	0	0
Dead B4	0	0	0	0	0	0	5	0	0	0	0
Dead B5	0	0	0	0	0	0	0	5	0	0	0
Dead B8	0	0	0	0	0	0	0	0	5	0	0
Dead B9	0	0	1	0	2	0	0	2	0	0	0
Medium	0	0	0	0	0	0	0	0	0	0	10

85% Correct, 87% Variance (7PCs)

B10: *Acinetobacterium*

B5: *E. coli*

B9: *Klebsiella pneumoniae*

B4: *M. luteus*

B8: *Salmonella*

Acknowledgements

ONR



DARPA



DOE



NIH



NSF



Walt Group

Myoyong Lee
Karri Michael
Paul Pantano
Caroline Schauer
Jenny Tam

Keith Albert
Todd Dickinson
Jane Ferguson
Michael Fleming
Jason Epstein

Ferenc Szurdoki
Frank Steemers
Shannon Stitzel
Laura Taylor
Tarun Mandal

Lawrence Livermore

Fred Milanovich

Pennsylvania State University

Peter C. Jurs

Tufts University School of Medicine

John S. Kauer Joel White

Johns Hopkins University

Lenore Cowen

Illumina

Mark Chee Kevin Gunderson